

Set Items Description
 S1 26174 SOMATOTROPIN
 S2 46607 GROWTH()HORMONE
 S3 57585 S1 OR S2
 S4 118932 MUTATION
 S5 266 S3 AND S4
 S6 22392 HELIX
 S7 6 S5 AND S6
 S8 3 AU="LEBENS M R" OR AU="LEBENS MR"
 S9 1 AU="TARDY D"
 S10 1000 AU="FISCHER M"
 S11 0 S3 AND S10
 S12 1230 E3-E37
 S13 0 S3 AND S12
 S14 0 S1 AND S12
 S15 19693 TUMOR()NECROSIS()FACTOR
 S16 299981 SEQUENCE
 S17 839 S15 AND S16
 S18 617 RD (unique items)
 S19 342669 RECEPTOR
 S20 1994 S15 AND S19
 S21 110839 SOLUBLE
 S22 233 S20 AND S21
 S23 151 RD (unique items)
 S24 12 SEQUENCE
 S25 299981 SEQUENCE
 S26 14 S23 AND S25
 ?t s26/3/1-14

625668

7-10 92

26/3/1 (Item 1 from file: 155)

08072529 92210529

Identification of cysteine-rich domains of the type 1 tumor necrosis factor receptor involved in ligand binding.

Marsters SA; Frutkin AD; Simpson NJ; Fendly BM; Ashkenazi A

Department of Immunobiology, Genentech, Inc., South San Francisco, California 94080.

J Biol Chem (UNITED STATES) Mar 25 1992, 267 (9) p5747-50, ISSN 0021-9258 Journal Code: HIV

Languages: ENGLISH

Document type: JOURNAL ARTICLE

26/3/2 (Item 2 from file: 155)

08054029 92192029

Cytoplasmic truncation of the p55 tumour necrosis factor (TNF) receptor abolishes signalling, but not induced shedding of the receptor.

Brakebusch C; Nophar Y; Kemper O; Engelmann H; Wallach D

Department of Molecular Genetics and Virology, Weizmann Institute of Science, Rehovot, Israel.

EMBO J (ENGLAND) Mar 1992, 11 (3) p943-50, ISSN 0261-4189

Journal Code: EMB

Languages: ENGLISH

Document type: JOURNAL ARTICLE

26/3/3 (Item 3 from file: 155)

07997196 92135196

Characterization of a recombinant extracellular domain of the type 1 tumor necrosis factor receptor: evidence for tumor necrosis factor-alpha induced receptor aggregation.

Pennica D; Kohr WJ; Fendly BM; Shire SJ; Raab HE; Borchardt PE; Lewis M; Goeddel DV

Department of Molecular Biology, Genentech, Inc., South San Francisco,

NET ITEMS Index-term
E1 6 AU=WALLACH C B
E2 5 AU=WALLACH CB
E3 371 *AU=WALLACH D
E4 1 AU=WALLACH D E
E5 1 AU=WALLACH D F
E6 188 AU=WALLACH D F H
E7 2 AU=WALLACH D H
E8 2 AU=WALLACH D H F
E9 9 AU=WALLACH D P
E10 1 AU=WALLACH DE
E11 155 AU=WALLACH DF
E12 2 AU=WALLACH DPH

9/25/68

Enter P or E for more
?s e3

530 371 AU="WALL C D"
?s s15 and s30

15293 S.5
371 S30
S31 88 S15 AND ?
?s s20 and s30

1994 S20
371 S30
S32 33 S20 AND S30
?t s32/3/1-33

32/3/1 (Item 1 from file: 155)
08130484 92268484

Selective decrease in cell surface expression and mRNA level of the 55-kDa tumor necrosis factor receptor during differentiation of HL-60 cells into macrophage-like but not granulocyte-like cells.

Winzen R; Wallach D; Engelmann H; Nophar Y; Brakebusch C; Kemper O; Resch K; Holtmann H

Institute of Molecular Pharmacology, Medical School, Hannover, FRG.
J Immunol (UNITED STATES) Jun 1 1992, 148 (11) p3454-60, ISSN 0022-1767 Journal Code: IFB

Languages: ENGLISH

Document type: JOURNAL ARTICLE

32/3/2 (Item 2 from file: 155)
28054029 92192029

Cytoplasmic truncation of the p55 tumour necrosis factor (TNF) receptor abolishes signalling, but not induced shedding of the receptor.

Brakebusch C; Nophar Y; Kemper O; Engelmann H; Wallach D

Department of Molecular Genetics and Virology, Weizmann Institute of Science, Rehovot, Israel.

EMBO J (ENGLAND) Mar 1992, 11 (3) p943-50, ISSN 0261-4189
Journal Code: EMB

Languages: ENGLISH

Document type: JOURNAL ARTICLE

32/3/3 (Item 3 from file: 155)

18004967 92142967

Soluble and cell surface receptors for tumor necrosis factor.

Wallach D; Engelmann H; Nophar Y; Aderka D; Kemper O; Hornik V; Holtmann H; Brakebusch C

Department of Molecular Genetics and Virology, Weizmann Institute of Science, Rehovot, Israel.

Agents Actions Suppl (SWITZERLAND) 1991, 35 p51-7, ISSN 0379-0363
Journal Code: ZYH

Languages: ENGLISH

Document type: JOURNAL ARTICLE; REVIEW; REVIEW LITERATURE

32/3/4 (Item 4 from file: 155)

7983813 92121013

Stabilization of the bioactivity of tumor necrosis factor by its soluble receptors.

Aderka D; Engelmann H; Maor Y; Brakebusch C; Wallach D

Department of Medicine T, Tel-Aviv Medical Center, Sackler Faculty of Medicine, Tel-Aviv University, Israel.

J Exp Med (UNITED STATES) Feb 1 1992, 175 (2) p323-9, ISSN 0022-1007
Journal Code: IEM

Languages: ENGLISH

Document type: JOURNAL ARTICLE

32/3/5 (Item 5 from file: 155)

1873651 92011651

Human neutrophil elastase releases a ligand-binding fragment from the 55-kDa tumor necrosis factor (TNF) receptor. Comparison with the proteolytic activity responsible for shedding of TNF receptors from stimulated neutrophils.

094417 BIOSIS Number: 4E1E917

CELL SURFACE AND SOLUBLE TNF RECEPTORS

WALLACH D; ADERKA D; ENGELMANN H; NOPHAR Y; KEMPER O; HOLTMANN H;

BRAKEBUSCH C; VILLA S; GONDI F G; BUCCIARELLI U

DEP. MOL. GENET. VIROL., WEIZMANN INST. SCI., REHOVOT 76100, ISR.

OSAWA, T. AND B. BONAVIDA (ED.). TUMOR NECROSIS FACTOR:

STRUCTURE-FUNCTION RELATIONSHIP AND CLINICAL APPLICATION; 3RD INTERNATIONAL

CONFERENCE ON TUMOR NECROSIS FACTOR AND RELATED CYTOKINES, MAKUHARI, JAPAN,

NOVEMBER 21-25, 1990. IX+291P. S. KARGER AG: BASEL, SWITZERLAND; NEW YORK,

NEW YORK, USA. ILLUS. ISBN 3-8055-5458-3. 0 (0). 1992. 47-57. CODEN:

36660

Language: ENGLISH

Document Type: CONFERENCE PAPER

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Journal Code: PV3

Languages: ENGLISH

Document type: JOURNAL ARTICLE

26/3/9 (Item 9 from file: 155)

07526991 91045991

A second tumor necrosis factor receptor gene product can shed a naturally occurring tumor necrosis factor inhibitor.

Kohno T; Brewer MT; Baker SL; Schwartz PE; King MW; Hale KK; Squires CH; Thompson RC; Vannice JL

Synergen, Inc., Boulder, CO 80301.

Proc Natl Acad Sci U S A Nov 1990, 87 (21) p8331-5, ISSN 0027-8424

Journal Code: PV3

Languages: ENGLISH

Document type: JOURNAL ARTICLE

26/3/10 (Item 10 from file: 155)

07498509 91017509

Cloning of human tumor necrosis factor (TNF) receptor cDNA and expression of recombinant soluble TNF-binding protein.

Gray PW; Barrett K; Chantry D; Turner M; Feldmann M

Charing Cross Sunley Research Centre, Hammersmith, London, England.

Proc Natl Acad Sci U S A Oct 1990, 87 (19) p7380-4, ISSN 0027-8424

Journal Code: PV3

Languages: ENGLISH

Document type: JOURNAL ARTICLE

26/3/11 (Item 11 from file: 155)

07487021 91006021

Soluble forms of tumor necrosis factor receptors (TNF-Rs). The cDNA for the type I TNF-R, cloned using amino acid sequence data of its soluble form, encodes both the cell surface and a soluble form of the receptor.

Nophar Y; Kemper O; Brakebusch C; Englemann H; Zwang R; Aderka D; Holtmann H; Wallach D

Department of Molecular Genetics and Virology, Weizmann Institute of Science, Rehovot, Israel.

EMBO J Oct 1990, 9 (10) p3269-78, ISSN 0261-4189 Journal Code: EMB

Languages: ENGLISH

Document type: JOURNAL ARTICLE

26/3/12 (Item 12 from file: 155)

07328285 90235285

Molecular cloning and expression of a receptor for human tumor necrosis factor.

Schall TJ; Lewis M; Koller KJ; Lee A; Rice GC; Wong GH; Gatanaga T; Granger GA; Lenz R; Raab H; et al

Department of Molecular Biology, Genentech, Inc., South San Francisco, California 94080.

Cell Apr 20 1990, 61 (2) p361-70, ISSN 0092-8674 Journal Code: CQ4

Languages: ENGLISH

Document type: JOURNAL ARTICLE

26/3/13 (Item 13 from file: 155)

07203215 90110215

Two tumor necrosis factor-binding proteins purified from human urine. Evidence for immunological cross-reactivity with cell surface tumor necrosis factor receptors.

Engelmann H; Novick D; Wallach D

Department of Molecular Genetics and Virology, Weizmann Institute of Science, Rehovot, Israel.

J Biol Chem Jan 25 1990, 265 (3) p1531-6, ISSN 0021-9258

Journal Code: HIV

Languages: ENGLISH

Document type: JOURNAL ARTICLE

26/3/14 (Item 1 from file: 5)

0006-2960 Journal Code: A0G
Languages: ENGLISH
Document type: JOURNAL ARTICLE

26/3/4 (Item 4 from file: 155)

07966250 92104250

Functional significance of human vascular smooth muscle cell-derived interleukin 1 in paracrine and autocrine regulation pathways.

Loppnow H; Libby P

Department of Biochemistry, Forschungsinstitut Borstel, Germany.

Exp Cell Res (UNITED STATES) Feb 1992, 198 (2) p283-90, ISSN 0014-4827 Journal Code: EPB

Contract/Grant No.: HL34636

Languages: ENGLISH

Document type: JOURNAL ARTICLE

26/3/5 (Item 5 from file: 155)

07940859 92078859

A tumor necrosis factor (TNF) receptor-IgG heavy chain chimeric protein as a bivalent antagonist of TNF activity.

Peppel K; Crawford D; Beutler B

Howard Hughes Medical Institute, University of Texas Southwestern Medical Center, Dallas, Texas 75235.

J Exp Med (UNITED STATES) Dec 1 1991, 174 (6) p1483-9, ISSN 0022-1007

Journal Code: I2V

Contract/Grant No.: P01-DK42582-01

Languages: ENGLISH

Document type: JOURNAL ARTICLE

26/3/6 (Item 6 from file: 155)

07688415 91207415

T2 open reading frame from the Shope fibroma virus encodes a soluble form of the TNF receptor.

Smith CA; Davis T; Wignall JM; Din WS; Farrah T; Upton C; McFadden G; Goodwin RG

Immunex Corporation, Seattle, WA 98101.

Biochem Biophys Res Commun Apr 15 1991, 176 (1) p335-42, ISSN 0006-291X Journal Code: 9Y8

Languages: ENGLISH

Document type: JOURNAL ARTICLE

26/3/7 (Item 7 from file: 155)

07571841 91090841

Molecular cloning and expression of human and rat tumor necrosis factor receptor chain (p60) and its soluble derivative, tumor necrosis factor-binding protein.

Himmler A; Maurer-Fogy I; Kronke M; Scheurich P; Pfizenmaier K; Lantz M; Olsson I; Hauptmann R; Stratowa C; Adolf GR

Ernst Boehringer Institut, Bender + Co GesmbH, Vienna, Austria.

DNA Cell Biol Dec 1990, 9 (10) p705-15, ISSN 0044-5498

Journal Code: AF9

Languages: ENGLISH

Document type: JOURNAL ARTICLE

26/3/8 (Item 8 from file: 155)

07543364 91062364

Purification and characterization of an inhibitor (soluble tumor necrosis factor receptor) for tumor necrosis factor and lymphotoxin obtained from the serum ultrafiltrates of human cancer patients.

Gatanaga T; Hwang CD; Kohr W; Cappuccini F; Lucci JA 3d; Jeffes EW; Lentz R; Tomich J; Yamamoto RS; Granger GA

Department of Molecular Biology and Biochemistry, University of California, Irvine 92717.

Proc Natl Acad Sci U S A Nov 1990, 87 (22) p8781-4, ISSN 0027-8424

(FILE 'USPAT' ENTERED AT 10:58:14 ON 10 JUL 92)

DEL HIS

L1 254 S TUMOR NECROSIS FACTOR
L2 201937 S SEQUENCE
L3 116 S L1 AND L2

=> d 13 cit 400-116

100. 4,785,077, Nov. 15, 1988, Substantially pure cytotoxicity triggering factor; Richard Kornbluth, et al., 530/351; 424/85.1, 85.4; 435/70.3, 70.4, 240.31, 948; 530/350, 827

101. 4,777,242, Oct. 11, 1988, Purification of recombinant **tumor** **necrosis** **factor**; Lynn P. Nelles, 530/351; 435/69.5, 803; 530/412, 415, 416, 417, 820

102. 4,770,995, Sep. 13, 1988, Detection of the sensitivity of cells to the effects of **tumor** **necrosis** **factor** and **'mphotoxin; Berish Y. Rubin, et al., 435/7.20, 436/501, 544, 545, 546

103. 4,752,565, Jun. 21, 1988, Oxidation-resistant mureins; Kirsten E. Koths, et al., 435/252.33, 172.3, 252.3, 256, 320.1; 530/351; 536/27; 930/141, 142; 935/10, 111

104. 4,748,234, May 31, 1988, Process for recovering ^{1/2} 668 bodies containing heterologous proteins from microbial hosts; Glenn Dorin, et al., 530/412; 435/69.51, 69.52; 530/351, 414, 416, 417, 422, 424, 829 ²

105. 4,721,671, Jan. 26, 1988, Efficient prokaryotic expression system using portions of the E. coli .beta.; Algis Anilionis, et al., 435/69.7, 172.3, 201, 252.33, 320.1, 849; 530/350; 536/27; 930/10, 240; 935/14 [IMAGE AVAILABLE]

106. 4,684,623, Aug. 4, 1987, Use of **tumor** **necrosis** **factor** as a weight regulator; James W. Lerrick, et al., 514/12; 424/85.1, 85.8; 514/11, 13, 14; 530/389.2; 930/144

107. 4,677,197, Jun. 30, 1987, Purification method for **tumor** **necrosis** **factor**; Leo S. Lin, et al., 530/417, 351, 416; 930/144, DIG.530

108. 4,677,064, Jun. 30, 1987, Human **tumor** **necrosis** **factor**; David F. Mark, et al., 435/69.1; 424/85.1, 88; 435/172.3, 252.3, 320.1; 514/8; 530/350, 351, 395, 808; 536/27; 930/144

109. 4,677,063, Jun. 30, 1987, Human **tumor** **necrosis** **factor**; David F. Mark, et al., 435/69.5; 424/85.1, 88; 435/172.3, 240.2, 252.3, 252.33, 320.1; 514/8, 12; 530/350, 351, 395, 808; 536/27; 930/144

110. 4,656,132, Apr. 7, 1987, Method of improving the yield of heterologous protein produced by cultivating recombinant bacteria; Arie Ben-Bassat, et al., 435/69.5, 69.51, 69.52, 172.3; 935/33, 38, 111

111. 4,650,674, Mar. 17, 1987, Synergistic cytotoxic composition; Bharat B. Aggarwal, et al., 424/85.5, 85.4; 435/69.5; 514/12; 930/143, 144

112. 4,588,585, May 13, 1986, Human recombinant cysteine depleted interferon-.beta. mureins; David F. Mark, et al., 424/85.2, 85.6; 435/69.51, 172.3; 530/351; 930/141, 142

113. 4,530,787, Jul. 23, 1985, Controlled oxidation of microbially produced cysteine-containing proteins; Ze'ev Shaked, et al., 530/351; 424/85.2, 85.4, 85.6, 88; 435/69.51, 811; 530/345, 363, 410, 808, 820, 825

114. 4,529,594, Jul. 16, 1985, Protein having antitumor activity; Hiroshi Hayashi, et al., 514/12; 424/85.1, 88; 530/324, 350, 413, 825, 828, 830; 930/10

115. 4,518,584, May 21, 1985, Human recombinant interleukin-2 mureins; David F. Mark, et al., 424/85.2, 85.1, 85.6; 435/69.51, 69.52, 172.3; 514/2, 8, 12; 530/351, 809; 930/141, 142

116. 4,481,137, Nov. 6, 1984, Glycoproteins and processes for their production; Haruo Ohnishi, et al., 530/395; 424/85.1; 435/70.3, 70.4; 514/8, 908; 530/322, 351, 806, 809, 828, 829

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(FILE 'USPAT' ENTERED AT 10:58:14 ON 10 JUL 92)

DEL HIS

L1 254 S TUMOR NECROSIS FACTOR
L2 201937 S SEQUENCE
L3 116 S L1 AND L2
L4 11032 S RECEPTOR
L5 82 S L1 AND L4

Porteu F; Brockhaus M; Wallach D; Engelmann H; Nathan L
Beatrice and Samuel A. Seaver Laboratory, Department of Medicine, Cornell
University Medical College, New York, New York 10021.

J Biol Chem Oct 5 1991, 266 (28) p18846-53, ISSN 0021-9258

Journal Code: HIV

Contract/Grant No.: CA45218

Languages: ENGLISH

Document type: JOURNAL ARTICLE

32/3/6 (Item 6 from file: 155)

07871844 92009844

The gene for the type II (p75) tumor necrosis factor receptor (TNF-RII) is localized on band 1p36.2-p36.3.

Kemper O; Derre J; Cherif D; Engelmann H; Wallach D; Berger R

Department of Molecular Genetics and Virology, Weitzmann Institute of Science, Rehovot, Israel.

Hum Genet Sep 1991, 87 (5) p623-4, ISSN 0340-6717 Journal Code: GED

Languages: ENGLISH

Document type: JOURNAL ARTICLE

32/3/7 (Item 7 from file: 155)

07867499 92005499

Increased serum levels of soluble receptors for tumor necrosis factor in cancer patients.

Aderka D; Englemann H; Hornik V; Skornick Y; Levo Y; Wallach D; Kushtai G

Department of Medicine, Tel-Aviv Sourasky Medical Center, Sackler School of Medicine, Tel Aviv University, Israel.

Cancer Res Oct 15 1991, 51 (20) p5602-7, ISSN 0008-5472

Journal Code: CNF

Languages: ENGLISH

Document type: JOURNAL ARTICLE

32/3/8 (Item 8 from file: 155)

07774800 91293800

The gene for the type I tumor necrosis factor receptor (TNF-R1) is localized on band 12p13.

Derre J; Kemper O; Cherif D; Nophar Y; Berger R; Wallach D

INSERM U301, Institut de Genetique Moleculaire, Paris, France.

Hum Genet Jun 1991, 87 (2) p231-3, ISSN 0340-6717 Journal Code: GED

Languages: ENGLISH

Document type: JOURNAL ARTICLE

32/3/9 (Item 9 from file: 155)

07766027 91285027

Induction of hyporesponsiveness to an early post-binding effect of tumor necrosis factor by tumor necrosis factor itself and interleukin 1.

Konig M; Wallach D; Resch K; Holtmann H

Institute of Molecular Pharmacology, Medical School, Hannover, FRG.

Eur J Immunol Jul 1991, 21 (7) p1741-5, ISSN 0014-2980

Journal Code: ENS

Languages: ENGLISH

Document type: JOURNAL ARTICLE

32/3/10 (Item 10 from file: 155)

07762541 91281541

Mechanisms controlling the level of receptors for tumor necrosis factor.

Holtmann H; Brakebusch C; Konig M; Klocke R; Winzen R; Resch K; Wallach D

Department of Molecular Pharmacology, Medical School Hannover, FRG.

Agents Actions Jan 1991, 32 (1-2) p106-8, ISSN 0065-4299

Journal Code: 2XZ

Languages: ENGLISH

Document type: JOURNAL ARTICLE

32/3/11 (Item 11 from file: 155)

07487021 91006021

Soluble forms of tumor necrosis factor receptors (TNF-Rs). The cDNA for

the type I TNF-R, cloned using amino acid sequence data or its soluble form, encodes both the cell surface and a soluble form of the receptor.

Nophar Y; Kemper O; Brakebusch C; Englemann H; Zwang R; Aderka D; Holtmann H; Wallach D
Department of Molecular Genetics and Virology, Weizmann Institute of Science, Rehovot, Israel.

EMBO J Oct 1990, 9 (10) p3269-78, ISSN 0261-4189 Journal Code: EMB

Languages: ENGLISH

Document type: JOURNAL ARTICLE

32/3/12 (Item 12 from file: 155)

07447443 90354443

Antibodies to a soluble form of a tumor necrosis factor (TNF) receptor have TNF-like activity.

Engelmann H; Holtmann H; Brakebusch C; Avni YS; Sarov I; Nophar Y; Hadas E; Leitner O; Wallach D
Department of Molecular Genetics and Virology, Weizmann Institute of Science, Rehovot, Israel.

J Biol Chem Aug 25 1990, 265 (24) p14497-504, ISSN 0021-9258

Journal Code: HIV

Languages: ENGLISH

Document type: JOURNAL ARTICLE

32/3/13 (Item 13 from file: 155)

07203215 90110215

Two tumor necrosis factor-binding proteins purified from human urine. Evidence for immunological cross-reactivity with cell surface tumor necrosis factor receptors.

Engelmann H; Novick D; Wallach D

Department of Molecular Genetics and Virology, Weizmann Institute of Science, Rehovot, Israel.

J Biol Chem Jan 25 1990, 265 (3) p1531-6, ISSN 0021-9258

Journal Code: HIV

Languages: ENGLISH

Document type: JOURNAL ARTICLE

32/3/14 (Item 14 from file: 155)

07006603 89308603

A tumor necrosis factor-binding protein purified to homogeneity from human urine protects cells from tumor necrosis factor toxicity.

Engelmann H; Aderka D; Rubinstein M; Rotman D; Wallach D

Department of Virology, Weizmann Institute of Science, Rehovot, Israel.

J Biol Chem Jul 15 1989, 264 (20) p11974-80, ISSN 0021-9258

Journal Code: HIV

Languages: ENGLISH

Document type: JOURNAL ARTICLE

32/3/15 (Item 15 from file: 155)

06553989 88198989

Dominance of resistance to the cytocidal effect of tumor necrosis factor in heterokaryons formed by fusion of resistant and sensitive cells.

Nophar Y; Holtmann H; Ber R; Wallach D

Department of Virology, Weizmann Institute of Science, Rehovot, Israel.

J Immunol May 15 1988, 140 (10) p3456-60, ISSN 0022-1767

Journal Code: IFB

Languages: ENGLISH

Document type: JOURNAL ARTICLE

32/3/16 (Item 16 from file: 155)

06307632 87281632

Down regulation of the receptors for tumor necrosis factor by interleukin 1 and 4 beta-phorbol-12-myristate-13-acetate.

Holtmann H; Wallach D

J Immunol Aug 15 1987, 139 (4) p1161-7, ISSN 0022-1767

Journal Code: IFB

Languages: ENGLISH

KEMPER O; DERRE J; CHERIF D; ENGELMANN H; WALLACH D; BERGER R
 UNITE INSERM U. 301, INS. GENETIQUE MOLECULAIRE, 27 RUE JULIETTE DODU,
 F-75010 PARIS, FR.

HUM GENET 87 (5). 1991. 623-624. CODEN: HUGED

Full Journal Title: Human Genetics

Language: ENGLISH

32/3/23 (Item 6 from file: 5)

8599313 BIOSIS Number: 92064313

THE GENE FOR THE TYPE 1 TUMOR NECROSIS FACTOR RECEPTOR TNF-R1 IS
 LOCALIZED ON BAND 12P13

DERRE J; KEMPER O; CHERIF D; NOPHAR Y; BERGER R; WALLACH D

INSERM U301, SDI NO. 15954 CNRS, INST. GENETIQUE MOLECULAIRE, 27 RUE
 JULIETTE DODU, F-7510 PARIS, FR.

HUM GENET 87 (2). 1991. 231-233. CODEN: HUGED

Full Journal Title: Human Genetics

Language: ENGLISH

32/3/24 (Item 7 from file: 5)

8445425 BIOSIS Number: 41129425

STUDYING STRUCTURE-FUNCTION RELATIONSHIP IN THE HUMAN TYPE 1 P55
 TNF-RECEPTOR USING HETEROLOGOUS EXPRESSION IN MURINE CELLS

BRAKEBUSCH C; NOPHAR Y; KEMPER O; ENGELMANN H; GRUNEWALD M; KING A;
 WALLACH D

SCH. MED., HANNOVER, FRG.

TWENTY-EIGHTH NATIONAL MEETING OF THE SOCIETY FOR LEUKOCYTE BIOLOGY AND
 THE TWENTY-FIRST LEUKOCYTE CULTURE CONFERENCE, ASPEN, COLORADO, USA,
 SEPTEMBER 28-OCTOBER 1, 1991. J LEUKOCYTE BIOL 0 (SUPPL. 2). 1991. 98.
 CODEN: JLBIE

Language: ENGLISH

Document Type: CONFERENCE PAPER

32/3/25 (Item 8 from file: 5)

7766655 BIOSIS Number: 90134655

SOLUBLE FORMS OF TUMOR NECROSIS FACTOR RECEPTORS TNF-Rs THE COMPLEMENTARY
 DNA FOR THE TYPE I TNF-R CLONED USING AMINO ACID SEQUENCE DATA OF ITS
 SOLUBLE FORM ENCODES BOTH THE CELL SURFACE AND A SOLUBLE FORM OF THE
 RECEPTOR

NOPHAR Y; KEMPER O; BRAKEBUSCH C; ENGELMANN H; ZWANG R; ADERKA D;
 HOLTZMANN H; WALLACH D

DEP. MOL. GENETICS VIROL., WEIZMANN INST. SCI., REHOVOT, ISRAEL 76100.

EMBO (EUR MOL BIOL ORGAN) J 9 (10). 1990. 3269-3278. CODEN: EMJOD

Full Journal Title: EMBO (European Molecular Biology Organization)

Journal

Language: ENGLISH

32/3/26 (Item 9 from file: 5)

7743514 BIOSIS Number: 90111514

ANTIBODIES TO A SOLUBLE FORM OF A TUMOR NECROSIS FACTOR TNF RECEPTOR HAVE
 TNF-LIKE ACTIVITY

ENGELMANN H; HOLTZMANN H; BRAKEBUSCH C; AVNI Y S; SAROV I; NOPHAR Y; HADAS
 E; LEITNER O; WALLACH D

DEP. MOL. GENETICS VIROL., WEIZMANN INST., REHOVOT 76100, ISRAEL.

J BIOL CHEM 265 (24). 1990. 14497-14504. CODEN: JBCHA

Full Journal Title: Journal of Biological Chemistry

Language: ENGLISH

32/3/27 (Item 10 from file: 5)

7548787 BIOSIS Number: 39061394

RECEPTOR-RELATED ASPECTS OF THE MODE OF ACTION OF TUMOR NECROSIS FACTOR
 AND ITS MODULATION

HOLTZMANN H; BRAKEBUSCH C; ENGELMANN H; KOENIG M; KLOCKE R; NOPHAR Y;
 RESCH K; WALLACH D

DEP. MOL. PHARMACOL., MEDICAL SCH. HANNOVER, FRG.

31ST SPRING MEETING OF THE DEUTSCHE GESELLSCHAFT FUER PHARMAKOLOGIE UND

32/3/17 (Item 17 from file: 155)

05948378 86249378

Binding of human TNF-alpha to high-affinity cell surface receptors: effect of IFN.

Israel S; Hahn T; Holtmann H; Wallach D

Immunol Lett Apr 1986, 12 (4) p217-24, ISSN 0165-2478

Journal Code: GIH

Languages: ENGLISH

Document type: JOURNAL ARTICLE

32/3/18 (Item 1 from file: 5)

8904179 BIOSIS Number: 42129179

CELL SURFACE AND SOLUBLE TNF RECEPTORS

WALLACH D; ADERKA D; ENGELMANN H; NOPHAR Y; KEMPER O; HOLTmann H;

BRAKEBUSCH C; VILLA S; GONDI F G; BUCCIARELLI U

DEP. MOL. GENET. VIROL., WEIZMANN INST. SCI., REHOVOT 76100, ISR.

OSAWA, T. AND B. BONAVIDA (ED.). TUMOR NECROSIS FACTOR: STRUCTURE-FUNCTION RELATIONSHIP AND CLINICAL APPLICATION; 3RD INTERNATIONAL CONFERENCE ON TUMOR NECROSIS FACTOR AND RELATED CYTOKINES, MAKUHARI, JAPAN, NOVEMBER 21-25, 1990. IX+291P. S. KARGER AG: BASEL, SWITZERLAND; NEW YORK, NEW YORK, USA. ILLUS. ISBN 3-8055-5458-3. 0 (0). 1992. 47-57. CODEN: 36660

Language: ENGLISH

Document Type: CONFERENCE PAPER

32/3/19 (Item 2 from file: 5)

8897179 BIOSIS Number: 42122179

CHANGES IN THE EXPRESSION OF TUMOR NECROSIS FACTOR RECEPTOR DURING DIFFERENTIATION OF HL-60 CELLS

HOLTmann H; WINZEN R; BRAKEBUSCH C; NOPHAR Y; ENGELMANN H; RESCH K; WALLACH D

INST. MOLECULAR PHARMACOL., MED. SCH., HANNOVER, FRG.

KEYSTONE SYMPOSIUM ON CYTOKINES IN GROWTH AND DEVELOPMENT, TAOS, NEW MEXICO, USA, FEBRUARY 1-8, 1992. J CELL BIOCHEM SUPPL 0 (16 PART B). 1992.

293. CODEN: JCBSD

Language: ENGLISH

Document Type: CONFERENCE PAPER

32/3/20 (Item 3 from file: 5)

8825551 BIOSIS Number: 42050551

CONTROL OF THE CELLULAR RESPONSE TO TUMOR NECROSIS FACTOR TNF

HOLTmann H; KOENIG M; ENGELMANN H; WALLACH D; RESCH K

INST. MOL. PHARMACOL., MED. SCH., 3000 HANNOVER 61, GER.

XXIIND MEETING OF THE SOCIETY OF IMMUNOLOGY, LUEBECK-TRAVEMUENDE, GERMANY, OCTOBER 23-26, 1991. IMMUNOBIOLOGY 183 (3-4). 1991. 257-258.

CODEN: IMMND

Language: ENGLISH

Document Type: CONFERENCE PAPER

32/3/21 (Item 4 from file: 5)

8667789 BIOSIS Number: 92132789

HUMAN NEUTROPHIL ELASTASE RELEASES A LIGAND-BINDING FRAGMENT FROM THE 75-KDA TUMOR NECROSIS FACTOR TNF RECEPTOR COMPARISON WITH THE PROTEOLYTIC ACTIVITY RESPONSIBLE FOR SHEDDING OF TNF RECEPTORS FROM STIMULATED NEUTROPHILS

PORTEU F; BROCKHAUS M; WALLACH D; ENGELMANN H; NATHAN C F

CORNELL UNIV. MED. COLL., BOX 57, 1300 YORK AVE., NEW YORK, N.Y. 10021.

J BIOL CHEM 266 (28). 1991. 18846-18853. CODEN: JBCHA

Full Journal Title: Journal of Biological Chemistry

Language: ENGLISH

32/3/22 (Item 5 from file: 5)

8659323 BIOSIS Number: 92124323

THE GENE FOR THE TYPE II P75 TUMOR NECROSIS FACTOR RECEPTOR TNF-RII IS

Language: ENGLISH

?log y

10Jul92 10:39:01 User217744 Session D126.2

\$8.28 0.230 Hrs File155

\$4.68 39 Type(s) in Format 3

\$4.68 39 Types

\$12.96 Estimated cost File155

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\$75.75 101 Types

\$104.46 Estimated cost File5

OneSearch, 2 files, 0.550 Hrs File05

\$5.94 DIALNET

\$123.36 Estimated cost this search

\$123.31 Estimated total session cost 0.556 Hrs.

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DIALNET: call cleared by request

Enter Service:

Language: ENGLISH

Document Type: CONFERENCE PAPER

32/3/28 (Item 11 from file: 5)

7151362 BIOSIS Number: 88074107

A TUMOR NECROSIS FACTOR-BINDING PROTEIN PURIFIED TO HOMOGENEITY FROM HUMAN URINE PROTECTS CELLS FROM TUMOR NECROSIS FACTOR TOXICITY

ENGELMANN H; ADERKA D; RUBINSTEIN M; ROTMAN D; WALLACH D

DEP. VIROL., WEIZMANN INST. SCI., REHOVOT, ISRAEL.

J BIOL CHEM 264 (20). 1989. 11974-11980. CODEN: JBCHA

Full Journal Title: Journal of Biological Chemistry

Language: ENGLISH

32/3/29 (Item 12 from file: 5)

6766030 BIOSIS Number: 36096551

IL-1 AND IFN MODULATE TNF-RECEPTOR EXPRESSION IN HUMAN POLYMORPHONUCLEAR AND MONONUCLEAR LEUKOCYTES

HAHN T; ENGELMANN H; HOLTZMANN H; LANDAU Z; WALLACH D

PEDIATRIC RES. INST., KAPLAN HOSP., REHOVOT.

ANNUAL INTERNATIONAL SOCIETY FOR INTERFERON RESEARCH MEETING ON INTERFERONS AND CYTOKINES, KYOTO, JAPAN, NOVEMBER 14-18, 1988. J INTERFERON RES 8 (SUPPL. 1). 1988. S148. CODEN: JIRED

Language: ENGLISH

Document Type: CONFERENCE PAPER

32/3/30 (Item 13 from file: 5)

6562182 BIOSIS Number: 86028733

DOMINANCE OF RESISTANCE TO THE CYTOCIDAL EFFECT OF TUMOR NECROSIS FACTOR IN HETEROKARYONS FORMED BY FUSION OF RESISTANT AND SENSITIVE CELLS

NOPHAR Y; HOLTZMANN H; BER R; WALLACH D

DEP. VIROL., WEIZMANN INST. SCI., REHOVOT 76100, ISRAEL.

J IMMUNOL 140 (10). 1988. 3456-3460. CODEN: JOIMA

Full Journal Title: Journal of Immunology

Language: ENGLISH

32/3/31 (Item 14 from file: 5)

6102312 BIOSIS Number: 34104619

REGULATION OF CELL RESPONSE TO TUMOR NECROSIS FACTOR

WALLACH D; NOPHAR Y; ADERKA D; ISRAEL S; HAHN T; ENGELMANN H; HOLTZMANN H
WEIZMANN INST. SCI., REHOVOT 76100, ISRAEL.

INTERNATIONAL CONFERENCE ON TUMOR NECROSIS FACTOR AND RELATED CYTOTOXINS, HEIDELBERG, WEST GERMANY, SEPTEMBER 14-18, 1987. IMMUNOBIOLOGY 175 (1-2).

1987. 42. CODEN: IMMND

Language: ENGLISH

Document Type: CONFERENCE PAPER

32/3/32 (Item 15 from file: 5)

5977388 BIOSIS Number: 84109953

DOWN REGULATION OF THE RECEPTORS FOR TUMOR NECROSIS FACTOR BY INTERLEUKIN 1 AND 4-BETA PHORBOL-12-MYRISTATE-13-ACETATE

HOLTZMANN H; WALLACH D

DEP. VIROL., WEIZMANN INST. SCI., REHOVOT 76100, ISR.

J IMMUNOL 139 (4). 1987. 1161-1167. CODEN: JOIMA

Full Journal Title: Journal of Immunology

Language: ENGLISH

32/3/33 (Item 16 from file: 5)

5409320 BIOSIS Number: 82054123

BINDING OF HUMAN TUMOR NECROSIS FACTOR TO HIGH-AFFINITY CELL SURFACE RECEPTORS EFFECT OF INTERFERON

ISRAEL S; HAHN T; HOLTZMANN H; WALLACH D

DEP. VIROL., WEIZMANN INST. SCI., REHOVOT, ISR.

IMMUNOL LETT 12 (4). 1986. 217-224. CODEN: IMLED

**** 15 standard deviations above mean ****							
7. N71302	HSV-1 cB and surrounding regi	3871	17	759	15.15	0	
8. Q10613	**** 3 standard deviations above mean ****						
9. Q06331	Rianodin receptor gene.	15672	22	700	13.95	0	
10. Q05872	Human IL-2R beta chain.	4035	22	695	13.84	0	
11. N70938	Sequence encoding human inter	4035	22	693	13.80	0	
12. N82026	DNA fragment contg. streptomy	3830	23	691	13.76	0	
13. Q03665	Cloned p2.1 insert encodes hu	3640	26	690	13.74	0	
14. N70834	Sequence homologous to Drosop	2465	22	689	13.72	0	
15. Q11707	L-Tryptophan production.	7726	21	687	13.68	0	
16. Q13402	Immunoglobulin D-region heavy	14928	23	687	13.68	0	
17. Q11010	Human alpha-2 plasmin inhibit	2243	15	687	13.68	0	
18. N92325	Fibulin C.	2200	19	683	13.60	0	
19. N80944	Human alpha-2-plasmin inhibit	2249	15	683	13.60	0	
20. Q05831	Alpha-2 plasmin inhibitor pre	2189	15	681	13.56	0	
	Cardiac sodium channel gene.	7555	21	681	13.56	0	

1. US-07-625-668A-1 (1-2175)

012215 Type I TNF receptor.

ID Q12215 standard; DNA; 2176 BP.
 AC Q12215;
 DT 12-SEP-1991 (first entry)
 DE Type I TNF receptor.
 KW Tumour Necrosis Factor; TNF; binding protein; TBP-I; ss.
 OS Homo sapiens.
 FH Key Location/Qualifiers
 FT terminator 244..246
 FT /*tag= a
 FT /note= "in-frame termination codon"
 FT CDS 256..1620
 FT /*tag= a
 FT /product= type I TNF receptor
 FT sig_peptide 256..318
 FT /*tag= b
 FT mat_peptide 319..1620
 FT /*tag= c
 FT misc_RNA 319..864
 FT /*tag= d
 FT /label= soluble_domain
 FT /note= "may be 2 codons shorter or a few codons
 longer"
 FT misc_feature 376..414
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 FT repeat_unit 505..633
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 FT repeat_unit 634..756
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 FT /number= 3
 FT repeat_unit 757..858
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 2020 2030 2040 2050 2060 2070 2080

 2090 2100 2110 2120 2130 2140 2150
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 TAAAAA
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 TAAAAA
 2170 X

2. US-07-625-668A-1 (1-2175)

006285 Human Tumour Necrosis Factor-Receptor cDNA insert.

ID 006285 standard; DNA; 2141 BP.
 AC 006285;
 DT 29-JAN-1991 (first entry).
 DE Human Tumour Necrosis Factor-Receptor cDNA insert.
 KW Tumour necrosis factor binding protein; TNF-BP; TNF-receptor;
 KW lambdaTNF-R2; ratTNF-R8; ss.
 OS Homo sapiens.
 FH Key Location/Qualifiers
 FT CDS 213..1577
 FT /*tag= a
 FT /label=huTNF-R
 PN EP-393438-A.
 PD 24-OCT-1990.
 PF 06-APR-1990; 106624.
 PR 21-APR-1989; DE-913101.
 PR 21-JUN-1989; DE-920282.
 PA (BOEH) BOEHRINGER INGELHEIMINT.
 PI Hauptmann R, Himmller A, Maurer-Fogy I, Stratowa C;
 DR WPI; 90-321987/43.
 DR P-PSDB; R07451.
 PT DNA encoding TNF binding protein and TNF- receptor - used in
 PT tumour treatment and to understand mechanismsm to TNF action
 PS Disclosure; Fig 91(1-2); 51pp; German.
 CC ratTNF-R8 (Q06284) was used to screen the HS913T cDNA library.
 CC LambdaTNF-R2 encodes the complete human TNF-R2 and was used to
 CC construct a plasmid (pADTNF-R) expressing the product the same way
 CC as pADTNF-BP (see Q06282).
 CC See also Q06282-Q06285.
 SQ Sequence 2141 BP; 455 A; 633 C; 593 G; 460 T;

Initial Score = 1711 Optimized Score = 2087 Significance = 42.35
 Residue Identity = 98% Matches = 2097 Mismatches = 35
 Gaps = 2 Conservative Substitutions = 0

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 ACTGGAGCCTCAGTCCAGAGAAATTCTGAGAAAATTAAAGCAGAGAGGGAGAGATCACTGGGACCAGGC
 ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
 GAATTCTCTGGACTGAGGCTCCAGTTCTGGCCTTGGGTTCA-AGATCACTGGGACCAGGC
 X 10 20 30 40 50 60

110 120 130 140 150 160 170
 CGTGATCTCTATGCCCGAGTCTCAACCCCTCAACTGTCACCCCAAGGCACCTGGGACGTCCCTGGACAGACCGA
 |||||||
 CGTGATCTCTATGCCCGAGTCTCAACCCCTCAACTGTCACCCCAAGGCACCTGGGACGTCCCTGGACAGACCGA

1230 1240 1250 1260 1270 1280 1290
ATCCTTGCACAGCCCTCGCCTCCGACCCCATCCCCAACCCCTTCAGAAGTGGGAGGACAGCGCCCACAAG
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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1230 1240 1250 1260 1270 1280 1290

1300 1310 1320 1330 1340 1350 1360
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1370 1380 1390 1400 1410 1420 1430 1440
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1450 1460 1470 1480 1490 1500 1510
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1520 1530 1540 1550 1560 1570 1580
CTGCTGGGACCGGTGCTCCGCGACATGGACCTGCTGGGCTGCCTGGAGGACATCGAGGAGGCCTTGCGGC
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1520 1530 1540 1550 1560 1570 1580

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CCC GCC CCCCCTCCGCCCGCCAGTCTTCTCAGATGAGGCTGCG-CCCTGCGGGCAGCTCTAAGGACCGT
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CCTGCGAGATCGCCTTCAACCCACTTTTCTGGAAAGGAGGGTCTGCAGGGGCAAGCAGGAGCTAGC
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1730 1740 1750 1760 1770 1780 1790
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1730 1740 1750 1760 1770 1780 1790 1800

1800 1810 1820 1830 1840 1850 1860 1870
CGCTGTGCGCGCGAGAGAGAGGTGCGCCGTGGGCTCAAGAGCCTGAGTGGTGGTTGCGAGGATGAGGGACG
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1810 1820 1830 1840 1850 1860 1870

1880 1890 1900 1910 1920 1930 1940
CTATGCCTCATGCCGTTTGGTGTCTCACCAAGCAAGGCTGCTGGGGGCCCTGGTTCGTCCCTGAGCC
||||||||||||||||||||||||||||||||||||
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1880 1890 1900 1910 1920 1930 1940

1950 1960 1970 1980 1990 2000 2010
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1950 1960 1970 1980 1990 2000 2010

440 450 460 470 480 490 500
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510 520 530 540 550 560 570
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580 590 600 610 620 630 640
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650 660 670 680 690 700 710 720
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940 950 960 970 980 990 1000
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1090 1100 1110 1120 1130 1140 1150
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1160 1170 1180 1190 1200 1210 1220
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PR 13-DEC-1989; IL-092697.
PR 12-JUL-1990; IL-095064.
PA (YEDA) YEDA RES & DEV CO LTD.
PI Wallach D, Nophar Y, Kemper O, Engelmann H, Brakebusch C;
PI Aderka D;
DR WPI; 91-186774/26.
DR P-PSDB; R12550.
PT Recombinant tumour necrosis factor binding protein I - prep'd. by
PT transfecting eukaryotic cells with vector contg. deoxyribonucleic
PT acid encoding human type T TNF receptor or soluble domain
PS Disclosure; Fig 1(D); 30pp; English.
CC The Tumour Necrosis Factor Binding Protein I is the soluble form of
CC type I TNF-receptor and constitutes a fragment of the cell surface
CC form of this receptor, corresp. to its extracellular domain.
CC There is no characteristic poly(A) addition signal near the 3' end
CC of the cDNA. The sequence ACTAAA (tag m) may serve as an
CC alternative to this signal, but with low efficiency.
CC See also Q12212-15.
SQ Sequence 2176 BP; 475 A; 644 C; 602 G; 455 T;

Initial Score = 1800 Optimized Score = 2167 Significance = 43.99
Residue Identity = 99% Matches = 2172 Mismatches = 3
Gaps = 1 Conservative Substitutions = 0

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X 10 20 30 40 50 60 70

80 90 100 110 120 130 140
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150 160 170 180 190 200 210
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220 230 240 250 260 270 280
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220 230 240 250 260 270 280

290 300 310 320 330 340 350 360
CCGCTGGTGCTCCTGGAGCTGTTGGTGGAAATATACCCCTCAGGGGTTATTGGACTGGTCCCTCACCTAGGG
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290 300 310 320 330 340 350 360

370 380 390 400 410 420 430
GACAGGGAGAAGAGAGATAGTGTGTCCCCAAGGAAAATATCCACCCCTCAAAATAATTGATTTGCTGT
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GACAGGGAGAAGAGAGATAGTGTCCTGGAGCTGTTGGTGGAAATATACCCCTCAGGGGTTATTGGACTGGTCCCTCACCTAGGG
370 380 390 400 410 420 430

Number of residues: 15631576
 Number of sequences optimized: 4191

The scores below are sorted by optimized score.
 Significance is calculated based on optimized score.

A 100% identical sequence to the query sequence was not found.

The list of best scores is:

12-13/96

Sequence Name	Description	Length	Init.	Opt.	Score	Score	Sig.	Frame
**** 47 standard deviations above mean ****								
1. HUMTNFRB	Homo sapiens tumor necrosis f	2111	1713	2085	47.96	0		
2. HUMTNFR	Human tumor necrosis factor r	2087	2074	2077	47.75	0		
3. HUMTNFRC	Human tumor necrosis factor r	2112	1660	2074	47.67	0		
**** 38 standard deviations above mean ****								
4. HSTNFBP	Human tumor necrosis factor r	2050	1625	1718	38.49	0		
5. HUMTNFRP	Human tumor necrosis factor r	2050	1625	1718	38.49	0		
**** 24 standard deviations above mean ****								
6. MUSTNFRE5	Mouse mRNA for 55-kDa tumor n	2179	817	1193	24.94	0		
7. MUSP55R	Murine mRNA for p55 tumor nec	2063	795	1160	24.09	0		
8. MUSMTNFR1	Mouse tumor necrosis factor r	2048	775	1159	24.07	0		
**** 23 standard deviations above mean ****								
9. MUSTNFR2	Murine tumor necrosis factor	1956	812	1136	23.47	0		
**** 21 standard deviations above mean ****								
10. RATTNFR	Rat tumor necrosis factor rec	2130	793	1042	21.05	0		
**** 12 standard deviations above mean ****								
11. CHKMYB15	Chicken c-myb oncogene DNA fo	8200	28	706	12.38	0		
12. PSESDSAB11	Pseudomonas sp. (strain ATCC	3510	25	704	12.33	0		
13. XAADHLA	X.autotrophicus haloalkane de	3041	28	697	12.15	0		
14. DROHBG	Drosophila melanogaster hunch	7680	23	696	12.12	0		
15. HUMINT2	Human int-2 proto-oncogene.	11608	25	696	12.12	0		
16. HUMTGFB	Human transforming growth fac	2744	24	695	12.10	0		
17. HS1GLYB	xerpes simplex virus type 1 g	9756	27	695	12.10	0		
18. M21629	Figure 2. Nucleotide sequence	9756	27	695	12.10	0		
19. HUMLARR	Human mRNA for LCA-homolog. L	7702	25	694	12.07	0		
20. SUSAL1COL	Strongylocentrotus purpuratus	4245	23	694	12.07	0		

1. US-07-625-668A-1 (1-2175)

HUMTNFRB Homo sapiens tumor necrosis factor receptor mRNA.

LOCUS HUMTNFRB 2111 bp ss-mRNA PRI 14-NOV-1990
 DEFINITION Homo sapiens tumor necrosis factor receptor mRNA, complete cds.
 ACCESSION M58286 M33480
 KEYWORDS tumor necrosis factor receptor.
 SOURCE Human cell line HL60, cDNA to mRNA.
 ORGANISM Homo sapiens
 Eukaryota; Animalia; Metazoa; Chordata; Vertebrata; Mammalia;
 Theria; Eutheria; Primates; Haplorhini; Catarrhini; Hominidae.
 REFERENCE 1 (bases 1 to 2111)
 AUTHORS Loetscher,H., Pan,Y.-C.E., Lahm,H.-W., Gentz,R., Brockhaus,M.,
 Tabuchi,H. and Lesslauer,W.
 TITLE Molecular cloning and expression of the human 55 kd tumor necrosis
 factor receptor
 JOURNAL Cell 61, 351-359 (1990)
 STANDARD simple staff_review
 FEATURES Location/Qualifiers
 CDS 187..1554
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2. US-07-625-668A-1 (1-2175)

HUMTNFR Human tumor necrosis factor receptor mRNA, comple

LOCUS HUMTNFR 2087 bp ss-mRNA PRI 15-SEP-1990
 DEFINITION Human tumor necrosis factor receptor mRNA, complete cds.
 ACCESSION M33294
 KEYWORDS cell surface receptor; tumor necrosis factor receptor.
 SOURCE Human placenta, cDNA to mRNA.
 ORGANISM Homo sapiens
 Eukaryota; Animalia; Metazoa; Chordata; Vertebrata; Mammalia;
 Theria; Eutheria; Primates; Haplorhini; Catarrhini; Hominidae.
 REFERENCE 1 (sites)
 AUTHORS Schall,T.J., Lewis,M., Koller,K.J., Lee,A., Rice,G.C., Wong,G.H.,
 Gatanaga,T., Granger,G.A., Lentz,R., Raab,H., Kohr,W.J. and
 Goeddel,D.V.
 TITLE Molecular cloning and expression of a receptor for human tumor
 necrosis factor
 JOURNAL Cell (1990) In press
 STANDARD full staff_review
 REFERENCE 2 (bases 1 to 2087)
 AUTHORS Schall,T.J., Lewis,M., Koller,K.J., Lee,A., Rice,G.C., Wong,G.H.,
 Gatanaga,T., Granger,G.A., Lentz,R., Raab,H., Kohr,W.J. and
 Goeddel,D.V.
 TITLE Molecular cloning and expression of a receptor for human tumor
 necrosis factor
 JOURNAL Cell 61, 361-370 (1990)
 STANDARD full staff_review
 COMMENT Draft entry and computer-readable sequence for [1] kindly submitted
 by T.Schall, 26-MAR-1990.
 FEATURES Location/Qualifiers
 CDS 182..1549
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3. US-07-625-668A-1 (1-2175)

HUMTNFRC Human tumor necrosis factor receptor (TNF receptor)

LOCUS HUMTNFRC 2112 bp ss-mRNA PRI 30-SEP-1991
 DEFINITION Human tumor necrosis factor receptor (TNF receptor) mRNA, complete
 cds.
 ACCESSION M63121 M75861
 KEYWORDS tumor necrosis factor receptor.
 SOURCE Human cDNA to mRNA.
 ORGANISM Homo sapiens
 Eukaryota; Animalia; Metazoa; Chordata; Vertebrata; Mammalia;
 Theria; Eutheria; Primates; Haplorhini; Catarrhini; Hominidae.
 REFERENCE 1 (bases 1 to 2112)
 AUTHORS Himmeler,A., Maurer-Fogy,I., Kroenke,M., Scheurich,P.,
 Pfizenmaier,K., Lantz,M., Olsson,I., Hauptmann,R., Stratowa,C. and
 Adolf,G.R.
 TITLE Molecular cloning and expression of human and rat tumor necrosis
 factor receptor chain (p60) and its soluble derivative, tumor
 necrosis factor-binding protein
 JOURNAL DNA Cell Biol. 9, 705-715 (1990)
 STANDARD simple staff entry
 FEATURES Location/Qualifiers
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500 510 520 530 540 550 560
620 630 640 650 660 670 680
CCGGGACACCGTGTGGCTGCAGGAAGAACAGTACCGCATTATTGGAGTGAACCTTTCCAGTGCTT
570 580 590 600 610 620 630
690 700 710 720 730 740 750
CAATTGCAGCCTCTGCCTCAATGGGACCGTGCACCTCTCCTGCCAGGAGAACAGAACACCGTGTGCACCTG
CAATTGCAGCCTCTGCCTCAATGGGACCGTGCACCTCTCCTGCCAGGAGAACAGAACACCGTGTGCACCTG
640 650 660 670 680 690 700
760 770 780 790 800 810 820 830
CCATGCAGGTTCTTCTAAGAGAAAACGAGTGTCTCCTGTAGTAACGTAAAGAAAAGCCTGGAGTGCAC
CCATGCAGGTTCTTCTAAGAGAAAACGAGTGTCTCCTGTAGTAACGTAAAGAAAAGCCTGGAGTGCAC
710 720 730 740 750 760 770 780
840 850 860 870 880 890 900
GAAGTTGTGCCTACCCAGATTGAGAATTTAAGGGCACTGAGGACTCAGGCACCAAGTGCTTGCCT
GAAGTTGTGCCTACCCAGATTGAGAATTTAAGGGCACTGAGGACTCAGGCACCAAGTGCTTGCCT
790 800 810 820 830 840 850
910 920 930 940 950 960 970
GGTCATTTCTTGGCTTGCCTTATCCCTCCTTCATTGGTTAATGTATCGCTACCAACGGTGGAA
GGTCATTTCTTGGCTTGCCTTATCCCTCCTTCATTGGTTAATGTATCGCTACCAACGGTGGAA
860 870 880 890 900 910 920
980 990 1000 1010 1020 1030 1040
GTCCAAGCTCTACTCCATTGTTGTGGAAATCGACACCTGAAAAAGAGGGGGAGCTTGAAGGAACACTAC
GTCCAAGCTCTACTCCATTGTTGTGGAAATCGACACCTGAAAAAGAGGGGGAGCTTGAAGGAACACTAC
930 940 950 960 970 980 990
1050 1060 1070 1080 1090 1100 1110
TAAGCCCCCTGGCCCCAAACCAAGCTTCAGTCCACTCCAGGCTTACCCCCACCCCTGGCTTCAGTCCCGT
TAAGCCCCCTGGCCCCAAACCAAGCTTCAGTCCACTCCAGGCTTACCCCCACCCCTGGCTTCAGTCCCGT
1000 1010 1020 1030 1040 1050 1060
1120 1130 1140 1150 1160 1170 1180 1190
GCCAGTTCCACCTTCACCTCCAGCTCCACCTATACCCCCGGTGACTGTCCCAACTTGCCTGGCTCCCCGAG
GCCAGTTCCACCTTCACCTCCAGCTCCACCTATACCCCCGGTGACTGTCCCAACTTGCCTGGCTCCCCGAG
1070 1080 1090 1100 1110 1120 1130 1140
1200 1210 1220 1230 1240 1250 1260
AGAGGTGGCACCCACCTATCAGGGGGCTGACCCCATCCTTGCAGACAGCCCTCGCCTCCGACCCCATCCCCAA
AGAGGTGGCACCCACCTATCAGGGGGCTGACCCCATCCTTGCAGACAGCCCTCGCCTCCGACCCCATCCCCAA
1150 1160 1170 1180 1190 1200 1210
1270 1280 1290 1300 1310 1320 1330
CCCCCTTCAGAAAGTGGAGGACAGCGCCACAAGCCACAGAGCCTAGACACTGATGACCCCGCGACGCTGTA
CCCCCTTCAGAAAGTGGAGGACAGCGCCACAAGCCACAGAGCCTAGACACTGATGACCCCGCGACGCTGTA
1220 1230 1240 1250 1260 1270 1280

4. US-07-625-668A-1 (1-2175)

Q10883 30kD TNF inhibitor precursor gene in lambda-gt10-7

ID Q10883 standard; cDNA; 2088 BP.
AC Q10883;
DT 13-MAY-1991 (first entry)
DE 30kD TNF inhibitor precursor gene in lambda-gt10-7ctnfbp.
KW Tumour necrosis factor; inhibitor; ss.
OS Homo sapiens.
FH Key Location/Qualifiers
FT CDS 171..1536
FT /*tag= a
PN AU9058976-A.
PD 24-JAN-1991.
PF 16-JUL-1990; 058976.
PR 18-JUL-1989; US-381080.
PR 11-DEC-1989; US-450329.
PR 07-FEB-1990; US-479661.
PA (SYNE-) SYNERGEN INC.
DR WPI; 91-073847/11.
DR P-PSDB; R10986.
PT Tumour necrosis factor inhibitor - for suppression of TNF-alpha
PT and -beta, useful as therapeutic agent.
PS Disclosure; Fig 21; 142pp; English.
CC The sequence encodes the entire 30 kD TNF inhibitor. The clone from
CC which the sequence was obtd. was isolated from a cDNA library
CC prep'd. from RNA form U937 cells treated with PMA/PHA. The whole
CC gene can be inserted into expression vectors for prepn. of TNF
CC inhibitor for use in the treatment of inflammatory and degenerative
CC diseases.
CC See also Q10878, Q10884 and Q10907.
SQ Sequence 2088 BP; 439 A; 626 C; 578 G; 445 T;

Initial Score = 1713 Optimized Score = 2077 Significance = 42.14
Residue Identity = 99% Matches = 2082 Mismatches = 5
Gaps = 1 Conservative Substitutions = 0

80 90 100 110 120 130 140
GGAGGGGAGAGATCACTGGGACCAAGGCCGTGATCTCTATGCCCGAGTCTCAACCCCTCAACTGTCACCCCAAG
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
GATCACTGGGACCAAGGCCGTGATCTCTATGCCCGAGTCTCAACCCCTCAACTGTCACCCCAAG
X 10 20 30 40 50 60

150 160 170 180 190 200 210 220
GCACCTGGGACGTCCTGGACAGACCGAGTCCCGGGAAAGCCCCAGCAGCACTGCCGCTGCCACACTGCCCTGAGCC
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
GCACCTGGGACGTCCTGGACAGACCGAGTCCCGGGAAAGCCCCAGCAGCACTGCCGCTGCCACACTGCCCTGAGCC
70 80 90 100 110 120 130

230 240 250 260 270 280 290
CAAATGGGGGAGTGAGAGGCCATAGCTGTCTGGCATGGGCCTCTCCACCGTGCCTGACCTGCTGCTGCCGCT
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
CAAATGGGGGAGTGAGAGGCCATAGCTGTCTGGCATGGGCCTCTCCACCGTGCCTGACCTGCTGCTGCCGCT
140 150 160 170 180 190 200

300 310 320 330 340 350 360
GGTGCTCTGGAGCTTTGGTGGGAATATAACCCCTCAGGGGTTATTGGACTGGTCCCTCACCTAGGGGACAG
||||||||||||||||||||||||||||||||||||||||||||||||||||
GGTGCTCTGGAGCTTTGGTGGGAATATAACCCCTCAGGGGTTATTGGACTGGTCCCTCACCTAGGGGACAG
210 220 230 240 250 260 270

370 380 390 400 410 420 430
GGAGAAAGAGAGATAGTGTGTGCCCCAAGGAAAATATCCACCCCTCAAAATAATTGATTTGCTGTACCAA
||||||||||||||||||||||||||||||||||||||||||||||||||||

GGAGAAGAGAGATAGTGTGTCCCCAAGGAAAATATATCCACCCCTCAAAATAATTGATTGCTGTACCAA
280 290 300 310 320 330 340 350
440 450 460 470 480 490 500
GTGCCACAAAGGAACCTACTTGTACAATGACTGTCCAGGCCGGGCAGGATACGGACTGCAGGGAGTGTGA
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
GTGCCACAAAGGAACCTACTTGTACAATGACTGTCCAGGCCGGGCAGGATACGGACTGCAGGGAGTGTGA
360 370 380 390 400 410 420
510 520 530 540 550 560 570 580
GAGCGGCTCCTCACCGCTTCAGAAAACCACCTCAGACACTGCCTCAGCTGCTCCAAATGCCGAAAGGAAAT
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
GAGCGGCTCCTCACCGCTTCAGAAAACCACCTCAGACACTGCCTCAGCTGCTCCAAATGCCGAAAGGAAAT
430 440 450 460 470 480 490
590 600 610 620 630 640 650
GGGTCAAGGTGGAGATCTCTTCTTGCACAGTGGACCGGGACACCGTGTGTGGCTGCAGGAAGAACCAAGTACCG
||||||||||||||||||||||||||||||||||||||||||||||||||||||||
GGGTCAAGGTGGAGATCTCTTCTTGCACAGTGGACCGGGACACCGTGTGTGGCTGCAGGAAGAACCAAGTACCG
500 510 520 530 540 550 560
660 670 680 690 700 710 720
GCATTATTGGAGTGAAAACCTTTCCAGTGCTTCATTGCAGCCTGCCTCAATGGGACCGTGCACCTCTC
||||||||||||||||||||||||||||||||||||||||||||||||
GCATTATTGGAGTGAAAACCTTTCCAGTGCTTCATTGCAGCCTGCCTCAATGGGACCGTGCACCTCTC
570 580 590 600 610 620 630
730 740 750 760 770 780 790
CTGCCAGGAGAAACAGAACACCGTGTGCACCTGCCATGCAGGTTCTTAAGAGAAAACGAGTGTGTCTC
||||||||||||||||||||||||||||||||||||||||||||
CTGCCAGGAGAAACAGAACACCGTGTGCACCTGCCATGCAGGTTCTTAAGAGAAAACGAGTGTGTCTC
640 650 660 670 680 690 700 710
800 810 820 830 840 850 860
CTGTAGTAACTGTAAGAAAAGCCTGGAGTGCACGAAGTTGTGCCTACCCAGATTGAGAATGTTAAGGGCAC
||||||||||||||||||||||||||||||||||||||||||||
CTGTAGTAACTGTAAGAAAAGCCTGGAGTGCACGAAGTTGTGCCTACCCAGATTGAGAATGTTAAGGGCAC
720 730 740 750 760 770 780
870 880 890 900 910 920 930 940
TGAGGACTCAGGCACCAAGTGTGTTGCCCTGGTCATTTCTTGGTCTTGCCTTTATCCCTCCTCTT
||||||||||||||||||||||||||||||||||||||||
TGAGGACTCAGGCACCAAGTGTGTTGCCCTGGTCATTTCTTGGTCTTGCCTTTATCCCTCCTCTT
790 800 810 820 830 840 850
950 960 970 980 990 1000 1010
CATTGGTTAATGTATCGTACCAACGGTGGAAAGTCAAGCTCTACTCCATTGTTGTGGAAATCGACACC
||||||||||||||||||||||||||||||||||||
CATTGGTTAATGTATCGTACCAACGGTGGAAAGTCAAGCTCTACTCCATTGTTGTGGAAATCGACACC
860 870 880 890 900 910 920
1020 1030 1040 1050 1060 1070 1080
TGAAAAAGAGGGGGAGCTTGAAGGAACTACTACTAACGCCCTGGCCCCAACCCAAGCTTCAGTCCCACCTCC
||||||||||||||||||||||||||||||||||||
TGAAAAAGAGGGGGAGCTTGAAGGAACTACTACTAACGCCCTGGCCCCAACCCAAGCTTCAGTCCCACCTCC
930 940 950 960 970 980 990
1090 1100 1110 1120 1130 1140 1150
AGGCTTCACCCCCCACCTGGCTTCAGTCCCACCTCCAGCTCCACCTTACACCTCCAGCTCCACCTATACCCC
||||||||||||||||||||||||||||||||
AGGCTTCACCCCCCACCTGGCTTCAGTCCCACCTCCAGCTCCACCTCCAGCTCCACCTATACCCC
1000 1010 1020 1030 1040 1050 1060 1070
1160 1170 1180 1190 1200 1210 1220
CGGTGACTGTCCCAACTTGCAGGCTCCCCGAGAGAGGTGGCACCAACCTATCAGGGGGCTGACCCATCCT
||||||||||||||||||||||||||||||||||||

CGGTGACTGTCCCAACTTGC GGCTCCCCCAGAGAGGTGGCACCCCTATCAGGGGGCTGACCCCATCCT
1080 1090 1100 1110 1120 1130 1140

1230 1240 1250 1260 1270 1280 1290 1300
TGCACAGCCCTGCCCTCCGACCCATCCCCAACCCCTTCAGAAGTGGGAGGACAGCGCCACAAGCCACA
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
TGCACAGCCCTGCCCTCCGACCCATCCCCAACCCCTTCAGAAGTGGGAGGACAGCGCCACAAGCCACA
1150 1160 1170 1180 1190 1200 1210

1310 1320 1330 1340 1350 1360 1370
GAGCCTAGACACTGATGACCCCCCGGACGCTGTACGCCGTGGAGAACGTGCCCCCGTTGCCCTGGAAGGA
||||||||||||||||||||||||||||||||||||||||||||||||||||||||
GAGCCTAGACACTGATGACCCCCCGGACGCTGTACGCCGTGGAGAACGTGCCCCCGTTGCCCTGGAAGGA
1220 1230 1240 1250 1260 1270 1280

1380 1390 1400 1410 1420 1430 1440
ATTCGTGCGCGCCTAGGGCTGAGCGACCAACGAGATCGATCGGCTGGAGCTGCAGAACGGCGCTGCCTGCG
||||||||||||||||||||||||||||||||||||||||||||||||||||
ATTCGTGCGCGCCTAGGGCTGAGCGACCAACGAGATCGATCGGCTGGAGCTGCAGAACGGCGCTGCCTGCG
1290 1300 1310 1320 1330 1340 1350

1450 1460 1470 1480 1490 1500 1510
CGAGGCGCAATACAGCATGCTGGCGACCTGGAGGGGGCGACGCCCGCGAGGCCACGCTGGAGCTGCT
||||||||||||||||||||||||||||||||||||||||||||||||
CGAGGCGCAATACAGCATGCTGGCGACCTGGAGGGGGCGACGCCCGCGAGGCCACGCTGGAGCTGCT
1360 1370 1380 1390 1400 1410 1420 1430

1520 1530 1540 1550 1560 1570 1580
GGGACCGCGTCTCCGCGACATGGACCTGCTGGAGGAGGACATCGAGGAGGCGCTTGC GGCCCCCGC
||||||||||||||||||||||||||||||||||||||||||||
GGGACCGCGTCTCCGCGACATGGACCTGCTGGAGGAGGACATCGAGGAGGCGCTTGC GGCCCCCGC
1440 1450 1460 1470 1480 1490 1500

1590 1600 1610 1620 1630 1640 1650 1660
CGCCCTCCGCCCGCGCCAGTCTTCTCAGATGAGGCTGCG-CCCTGCGGGCAGCTCTAAGGACCGTCCTGC
||||||||||||||||||||||||||||||||||||||||||||
CGCCCTCCGCCCGCGCCAGTCTTCTCAGATGAGGCTGCGCCCCCTGCGGGCAGCTCTAAGGACCGTCCTGC
1510 1520 1530 1540 1550 1560 1570

1670 1680 1690 1700 1710 1720 1730
GAGATCGCCTTCCAACCCCACTTTCTGAAAGGAGGGGTCTGCAGGGCAAGCAGGAGCTAGCAGCCG
||||||||||||||||||||||||||||||||||||||||
GAGATCGCCTTCCAACCCCACTTTCTGAAAGGAGGGGTCTGCAGGGCAAGCAGGAGCTAGCAGCCG
1580 1590 1600 1610 1620 1630 1640

1740 1750 1760 1770 1780 1790 1800
CCTACTTGGTGTCTAACCCCTCGATGTACATAGCTTCTCAGCTGCCTGCGCGCCGCCAGTCAGCGCTG
||||||||||||||||||||||||||||||||||||||||
CCTACTTGGTGTCTAACCCCTCGATGTACATAGCTTCTCAGCTGCCTGCGCGCCGCCAGTCAGCGCTG
1650 1660 1670 1680 1690 1700 1710

1810 1820 1830 1840 1850 1860 1870
TGC CGCGGGAGAGAGGGTGC CGCGGTGGGCTCAAGAGCCTGAGTGGGTTGCGAGGATGAGGGACGCTATG
||||||||||||||||||||||||||||||||||||||||
TGC CGCGGGAGAGAGGGTGC CGCGGTGGGCTCAAGAGCCTGAGTGGGTTGCGAGGATGAGGGACGCTATG
1720 1730 1740 1750 1760 1770 1780 1790

1880 1890 1900 1910 1920 1930 1940
CCTCATGCCGTTTGGGTGTCTCACCAAGCAAGGCTGCTCGGGGGCCCTGGTTCGTCCCTGAGCCTTTT
||||||||||||||||||||||||||||||||||||||||
CCTCATGCCGTTTGGGTGTCTCACCAAGCAAGGCTGCTCGGGGGCCCTGGTTCGTCCCTGAGCCTTTT
1800 1810 1820 1830 1840 1850 1860

1950 1960 1970 1980 1990 2000 2010 2020
CACAGTGCATAAGCAGTTTTTTGTGTTTGTGTTTGTGTTTAAATCAATCATGTTACACTAA
||||||||||||||||||||||||||||||||||||||||

CACAGTGCATAAGCAGTTTTTGTGTTTGTGTTTAAATCAATCATGTTACACTAA

1870 1880 1890 1900 1910 1920 1930

2030 2040 2050 2060 2070 2080 2090

TAGAAACTTGGCACTCCTGTGCCCTCTGCCCTGGACAAGCACATAGCAAGCTGAAC

TAGAAACTTGGCACTCCTGTGCCCTCTGCCCTGGACAAGCACATAGCAAGCTGAAC

1940 1950 1960 1970 1980 1990 2000

2100 2110 2120 2130 2140 2150 2160

GAGCACGGAACAAATGGGGCCTTCAGCTGGAGCTGGACTTTGTACATACACTAAAATTCTGAAGTTAAA

||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

GAGCACGGAACAAATGGGGCCTTCAGCTGGAGCTGGACTTTGTACATACACTAAAATTCTGAAGTTAAAG

2010 2020 2030 2040 2050 2060 2070

2170 X

AAAAAAA

||||||

CTCAAAA

2080 X

5. US-07-625-668A-1 (1-2175)

006282 Plasmid Tumour Necrosis Factor-Binding Protein 15

ID Q06282 standard; DNA; 1334 BP.

AC Q06282;

DT 29-JAN-1991 (first entry)

DE Plasmid Tumour Necrosis Factor-Binding Protein 15 cDNA insert.

KW Tumour necrosis factor binding protein; TNF-BP; TNF-receptor;

KW pTNF-BP15; ss.

DS Homo sapiens.

FH Key Location/Qualifiers

FT CDS 213..1325

FT /*tag= a

FT /product=TNF-BP

PN EP-393438-A.

PD 24-OCT-1990.

PF 06-APR-1990; 106624.

PR 21-APR-1989; DE-913101.

PR 21-JUN-1989; DE-920282.

PA (BOEH) BOEHRINGER INGELHEIMINT.

PI Hauptmann R, Himmller A, Maurer-Fogy I, Stratowa C;

DR WPI; 90-321987/43.

DR P-PSDB; R07449.

PT DNA encoding TNF binding protein and TNF- receptor - used in

PT tumour treatment and to understand mechanismsm to TNF action

PS Disclosure; Fig 1(1-3); 51pp; German.

CC pTNF-BP15 is one of 30 positives clones in a screened cDNA library

CC from induced TNF-induced fibrosarcoma cells. A TNF-BP had been

CC isolated from the urine of patients with uraemia and probes/primers

CC were constructed from the determined amino acid sequence.

CC To produce a vector expressing a soluble form of TNF-binding

CC protein, this plasmid was cut with XmmI, amplified by PCR and the

CC amplified DNA cut with BamHI and EcoRI.

CC The resulting 0.75 kb DNA fragment was inserted into pT7/T3 alpha-19

CC (BRL) cut with the same enzymes to recover pTNF-BP. This was cut

CC with BamHI and EcoRI, and the recovered fragment inserted into

CC pAD-CMV1 (Q06283) to give the required plasmid pADTNF-BP.

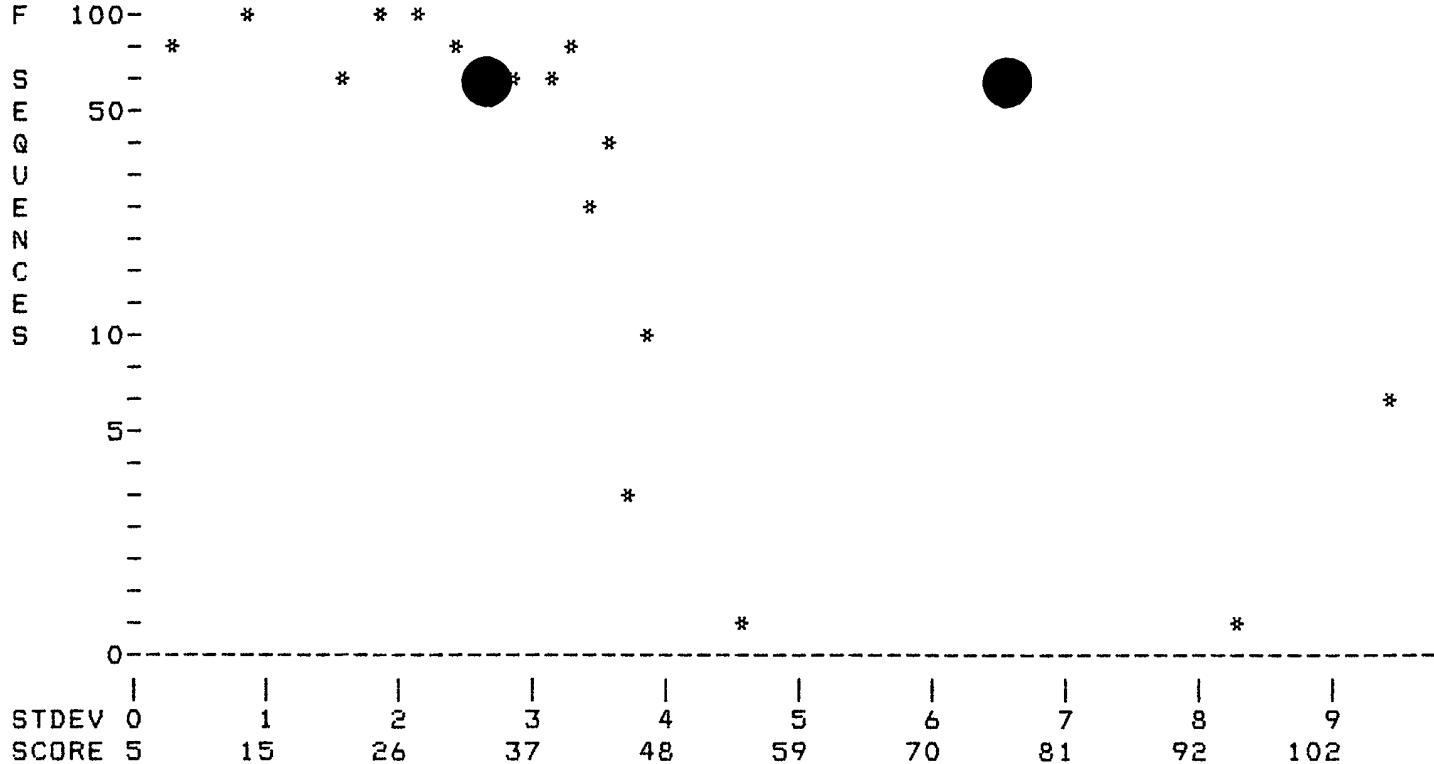
CC See also Q06282-Q06285.

SQ Sequence 1334 BP; 299 A; 409 C; 342 G; 284 T;

Initial Score = 1289 Optimized Score = 1298 Significance = 26.19

Residue Identity = 97% Matches = 1303 Mismatches = 31

bps = 1 Conservative Substitutions = 0



PARAMETERS

Similarity matrix	Unitary	K-tuple	2
Mismatch penalty	5	Joining penalty	20
Gap penalty	5.00	Window size	13
Gap size penalty	0.26		
Cutoff score	0		
Randomization group	0		
Initial scores to save	20	Alignments to save	10
Optimized scores to save	20	Display context	10

SEARCH STATISTICS

Scores:	Mean 5	Median 20	Standard Deviation 10.89
Times:	CPU 00:02:00.00	Total Elapsed 00:04:01.00	
Number of residues:	1428997		
Number of sequences optimized:	3856		

The scores below are sorted by optimized score.
Significance is calculated based on optimized score.

A 100% identical sequence to the query sequence was found:

Sequence Name	Description	Init.	Opt.	Length	Score	Score	Sig.	Frame
1. R10986	30KD TNF inhibitor precursor.	455	455	455	41.34	0		

The list of other best scores is:

Sequence Name	Description	Init.	Opt.	Length	Score	Score	Sig.	Frame
**** 41 standard deviations above mean ****								

2. R11082	Human 55kD TNF-binding protein	455	454	454	41.25	0
3. R12550	Type I TNF receptor.	455	453	453	41.15	0
	**** 40 standard deviations above mean ****					
4. R07451	Human Tumour Necrosis Factor-	455	451	451	40.97	0
	**** 33 standard deviations above mean ****					
5. R07449	Tumour Necrosis Factor-Bindin	371	371	371	33.62	0
	**** 19 standard deviations above mean ****					
6. R07450	Rat Tumour Necrosis Factor-Re	461	240	212	19.02	0
	**** 8 standard deviations above mean ****					
7. R10984	Partial sequence 30kD TNF inh	102	81	96	8.36	0
	**** 4 standard deviations above mean ****					
8. R11142	TNF-R deduced from mTNF-R clo	474	8	55	4.59	0
	**** 3 standard deviations above mean ****					
9. P70416	Polypeptide with IgE binding	557	5	47	3.86	0
10. R13949	SUP-B27 t(1;19) translocation	736	9	47	3.86	0
11. R12914	Human Thyroid Stimulating Hor	764	6	47	3.86	0
12. R13269	Human Thyroid Stimulating Hor	764	6	47	3.86	0
13. R12505	Human thyroid stimulating hor	764	6	47	3.86	0
14. R13951	E2A/pri fusion protein TYPE I	825	9	47	3.86	0
15. R15158	E2A/pri fusion protein TYPE I	742	9	47	3.86	0
16. R13948	SUP-B27 t(1;19) translocation	819	9	47	3.86	0
17. R10656	Hepatic parenchymal cell grow	727	8	46	3.77	0
18. P61048	A.nidulans phosphoenol pyruvic	1085	5	46	3.77	0
19. R11254	Human IL-4 receptor.	825	5	46	3.77	0
20. R08406	Sequence deduced from env gen	846	6	45	3.67	0

1. US-07-625-668A-2 (1-455)

R10986 30kD TNF inhibitor precursor.

ID R10986 standard; Protein: 455 AA.
 AC R10986;
 DT 13-MAY-1991 (first entry)
 DE 30kD TNF inhibitor precursor.
 KW Tumour necrosis factor; inhibitor.
 OS Homo sapiens.
 FH Key Location/Qualifiers
 FT Cleavage-site 40..41
 FT note=" cleavage gives active protein "
 PN AU9058976-A.
 PD 24-JAN-1991.
 PF 16-JUL-1990; 058976.
 PR 18-JUL-1989; US-381080.
 PR 11-DEC-1989; US-450329.
 PR 07-FEB-1990; US-479661.
 PA (SYNE-) SYNERGEN INC.
 DR WPI; 91-073847/11.
 DR N-PSDB; Q10883.
 PT Tumour necrosis factor inhibitor - for suppression of TNF-alpha
 PT and -beta, useful as therapeutic agent.
 PS Disclosure; Fig 21; 142pp; English.
 CC The sequence comprises the entire 30 kD TNF inhibitor. The clone
 CC from which the sequence was deduced was isolated from a cDNA
 CC library prep'd. from RNA form U937 cells treated with PMA/PHA.
 CC The whole gene can be inserted into expression vectors for prep'n.
 CC of TNF inhibitor for use in the treatment of inflammatory and
 CC degenerative diseases. The active protein is claimed (Claim 8).
 CC See also R10984 and R11001.
 SQ Sequence 455 AA;
 SQ 20 A; 27 R; 17 N; 21 D; 0 B; 30 C; 15 Q; 30 E; 0 Z; 30 G; 10 H;
 SQ 13 I; 58 L; 20 K; 5 M; 14 F; 37 P; 36 S; 31 T; 5 W; 13 Y; 23 V;
 Initial Score = 455 Optimized Score = 455 Significance = 41.34
 Residue Identity = 100 Matches = 455 Mismatches = 0
 Gaps = 0 Conservative Substitutions = 0

PD 20-MAR-1991.
PF 31-AUG-1990; 116707.
PR 12-SEP-1989; CH-003319.
PR 08-MAR-1990; CH-000746.
PR 20-APR-1990; CH-001347.
PA (HOFF) HOFFMANN-LA ROCHE AG.
PI Brockhaus M, Dembic Z, Gentz R, Lesslauer W, Lotscher H;
PI Schlaeger EJ;
DR WPI; 91-081851/12.
DR N-PSDB; Q10955.
PT Insoluble tumour necrosis factor binding proteins - and DNA
PT encoding them, useful in pharmaceutical prods. and for antibody
PT prodn.
PS Claim 1; Fig 1; 26pp; German.
CC Partial amino acid sequences were determined for the 55 and 75kD
CC TNF-BPs (see R11072-R11081) and oligonucleotide primers were
CC synthesised based on these partial sequences. The primers were used
CC to produce a cDNA fragment for use as a probe to screen a human
CC placental cDNA bank constructed in lambda gt11. Positive clones were
CC identified and sequenced. DNA constructs comprising the TNF-BP coding
CC sequence may also contain a fragment encoding a human Ig domain.
CC Recombinant constructs are used to transform cells to confer
CC improved TNF-binding properties.
CC See also Q10956.
SQ Sequence 455 AA;
SQ 20 A; 27 R; 17 N; 21 D; 0 B; 30 C; 15 Q; 30 E; 0 Z; 30 G; 10 H;
SQ 13 I; 57 L; 20 K; 5 M; 14 F; 37 P; 36 S; 31 T; 5 W; 13 Y; 24 V;

Initial Score = 454 Optimized Score = 454 Significance = 41.25
Residue Identity = 99% Matches = 454 Mismatches = 1
Gaps = 0 Conservative Substitutions = 0

X 10 20 30 40 50 60 70
MGLSTVPDLLLPLVLELLVGIYPSGVIGLVPHLGDREKRDSVCPQGKYIHPQNNSICCTKCHKGTYLYNDC
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
MGLSTVPDLLLPLVLELLVGIYPSGVIGLVPHLGDREKRDSVCPQGKYIHPQNNSICCTKCHKGTYLYNDC
X 10 20 30 40 50 60 70

80 90 100 110 120 130 140
PGPGQDTDCRECESGSFTASENHRLHCLSCSKCRKEMGQVEISSCTVDRDTVCGRKNQYRHYSNLFQCF
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
PGPGQDTDCRECESGSFTASENHRLHCLSCSKCRKEMGQVEISSCTVDRDTVCGRKNQYRHYSNLFQCF
80 90 100 110 120 130 140

150 160 170 180 190 200 210
NCSLCLNGTVHLSCQEKGNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
NCSLCLNGTVHLSCQEKGNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL
150 160 170 180 190 200 210

220 230 240 250 260 270 280
VIFFGLCLLSLLFIGLMYRYQRWKSCLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFSV
||||||||||||||||||||||||||||||||||||||||||||||||||||||||
VIFFGVCLLSLLFIGLMYRYQRWKSCLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFSV
220 230 240 250 260 270 280

290 300 310 320 330 340 350 360
PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQGADPILATALASDPIPNPLQKWEDSAHKPQSLDTDDPATLY
||||||||||||||||||||||||||||||||||||||||||||||||||||
PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQGADPILATALASDPIPNPLQKWEDSAHKPQSLDTDDPATLY
290 300 310 320 330 340 350 360

370 380 390 400 410 420 430
AVVENVPPLRWKEFVRLGL [REDACTED] HEIDRLELQNGRCLREAQYSMLATW [REDACTED] TPRREATLELLGRVLRDMDDLLG
||||||||||||||||||||||||||||||||||||||||||||||||
AVVENVPPLRWKEFVRLGLSDHEIDRLELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRDMDDLLG

X	10	20	30	40	50	60	70
MGLSTVPDLLLPLVLLVELLVGIYPSGVIGLVPHLGDREKRDSVCPQGKYIHPQNNNSICCTKCHKGTYLYND							
X	10	20	30	40	50	60	70
80	90	100	110	120	130	140	
PGPGQDTDCRECESGSFTASENHRLHCLSCSKCRKEMGQVEISSCTVDRDTVCGCRKNQYRHYWSENLFQCF							
PGPGQDTDCRECESGSFTASENHRLHCLSCSKCRKEMGQVEISSCTVDRDTVCGCRKNQYRHYWSENLFQCF							
80	90	100	110	120	130	140	
150	160	170	180	190	200	210	
NCSLCLNGTVHLSCQEKGQNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL							
NCSLCLNGTVHLSCQEKGQNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL							
150	160	170	180	190	200	210	
220	230	240	250	260	270	280	
VIFFGLCLLSLLFIGLMYRYQRWKSCLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFSPV							
VIFFGLCLLSLLFIGLMYRYQRWKSCLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFSPV							
220	230	240	250	260	270	280	
290	300	310	320	330	340	350	360
PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQGADPILATALASDPIPPLQKWEDESAHKPQSLDDPATLY							
PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQGADPILATALASDPIPPLQKWEDESAHKPQSLDDPATLY							
290	300	310	320	330	340	350	360
370	380	390	400	410	420	430	
AVVENVPPLRWKEFVRRGLSDHEIDRLELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRDM DLLG							
AVVENVPPLRWKEFVRRGLSDHEIDRLELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRDM DLLG							
370	380	390	400	410	420	430	
440	450	X					
CLEDIEEALCGPAALPPAPSLLR							
CLEDIEEALCGPAALPPAPSLLR							
440	450	X					

2. US-07-625-668A-2 (1-455)

R11082 Human 55kD TNF-binding protein.

ID R11082 standard; Protein; 455 AA.
 AC R11082;
 DT 24-MAY-1991 (first entry)
 DE Human 55kD TNF-binding protein.
 KW Tumour Necrosis Factor; binding proteins; septic shock;
 KW autoimmune glomerulonephritis; lymphokine; cytokine.
 FH Key Location/Qualifiers
 FT Modified -site 54
 FT /label= putative N-glycosylation site
 FT Modified -site 145
 FT /label= putative N-glycosylation site
 FT Modified -site 151
 FT /label= putative N-glycosylation site
 FT Modified -site 270
 FT /label= putative N-glycosylation site
 FT Region 212..230
 FT /label= transmembrane region
 FT Peptide 1..28
 FT /label= signal peptide
 PN EP-417563-A.

370

380

390

400

410

420

430

440 450 X
 CLEDIEEALCGPAALPPAPSLLR
 |||||||||||||||||||||
 CLEDIEEALCGPAALPPAPSLLR
 440 450 X

3. US-07-625-668A-2 (1-455)

R12550 Type I TNF receptor.

ID R12550 standard; Protein; 455 AA.
 AC R12550;
 DT 12-SEP-1991 (first entry)
 DE Type I TNF receptor.
 KW Tumour Necrosis Factor; TNF; binding protein; TBP-I.
 OS Homo sapiens.
 FH Key Location/Qualifiers
 FT Peptide 1..21
 FT /label= sig_peptide
 FT Protein 22..455
 FT /label= mat_protein
 FT Domain 212..234
 FT /label= transmembrane_domain
 FT Region 41..53
 FT /label= TBP-I derived sequence
 FT Region 110..124
 FT /label= TBP-I derived sequence
 FT Region 199..201
 FT /label= TBP-I derived sequence
 FT Modified -site 54..56
 FT /label= N-glycosylation_site
 FT Modified -site 145..147
 FT /label= N-glycosylation_site
 FT Modified -site 151..153
 FT /label= N-glycosylation_site
 FT Region 44..83
 FT /label= repeat_1
 FT Region 84..126
 FT /label= repeat_2
 FT Region 127..167
 FT /label= repeat_3
 FT Region 168..201
 FT /label= repeat_4
 FT Domain 21..203
 FT /label= soluble_domain
 FT /note= "may be 2 amino acids shorter or contain a
 few additional amino acids" 90
 PN EP-433900-A. 13
 PD 26-JUN-1991. 12
 PF 13-DEC-1990; 124133. 12
 PR 13-DEC-1989; IL-092697.
 PR 12-JUL-1990; IL-095064.
 PA (YEDA) YEDA RES & DEV CO LTD.
 PI Wallach D, Nophar Y, Kemper O, Engelmann H, Brakebusch C;
 PI Aderka D;
 DR WPI; 91-186774/26.
 DR N-PSDB; Q12215.
 PT Recombinant tumour necrosis factor binding protein I - prep'd. by
 PT transfecting eukaryotic cells with vector contg. deoxyribonucleic
 PT acid encoding human type I TNF receptor or soluble domain
 PS Disclosure; Fig 1(D); 30pp; English.
 CC The Tumour Necrosis Factor Binding Protein I is the soluble form of
 CC type I TNF-receptor and constitutes a fragment of the cell surface
 CC form of this receptor, corresp. to its extracellular domain.

CC The soluble proteins produced by the transfected cells secreted
CC into the medium may have at the N-terminus the sequence Asp-Ser-Val
CC (41-43), or the sequence Leu-Val-Pro (30-32) or Ile-Tyr-Pro (22-24)
CC or any other sequence between Ile (21) and Asp (41).

CC See also Q12212-15.

SQ Sequence 455 AA;

SQ 20 A; 27 R; 17 N; 21 D; 0 B; 30 C; 15 Q; 30 E; 0 Z; 30 G; 10 H;

SQ 13 I; 58 L; 20 K; 5 M; 14 F; 37 P; 37 S; 31 T; 4 W; 13 Y; 23 V;

Initial Score = 453 Optimized Score = 453 Significance = 41.15
Residue Identity = 99% Matches = 453 Mismatches = 2
Gaps = 0 Conservative Substitutions = 0

X 10 20 30 40 50 60 70
MGLSTVPDLLLPLVLELLVGIYPSGVIGLVPHLGDREKRDSVCPQGKYIHPQNNNSICCTKCHKGTYLYND
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
MGLSTVPDLLLPLVLELLVGIYPSGVIGLVPHLGDREKRDSVCPQGKYIHPQNNNSICCTKCHKGTYLYND
X 10 20 30 40 50 60 70

80 90 100 110 120 130 140
PGPGQDTDCRECESGSFTASENHLRHCLSCSKCRKEMGQVEISSCTVDRDTVCGRKRNQYRHYWSENLFQCF
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
PGPGQDTDCRECESGSFTASENHLRHCLSCSKCRKEMGQVEISSCTVDRDTVCGRKRNQYRHYWSENLFQCF
80 90 100 110 120 130 140

150 160 170 180 190 200 210
NCSLCLNGTVHLSCQEKKQNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
NCSLCLNGTVHLSCQEKKQNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL
150 160 170 180 190 200 210

220 230 240 250 260 270 280
VIFFGLCLLSLLFIGLMYRYQRWKSCLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFSPV
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
VIFFGLCLLSLLFIGLMYRYQRCKSKLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFSPV
220 230 240 250 260 270 280

290 300 310 320 330 340 350 360
PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQGADPILATALASDPIPPLQKEDSAHKPQSLDDPATLY
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQGADPILATALASDPIPPLQKEDSAHKPQSLDDPATLY
290 300 310 320 330 340 350 360

370 380 390 400 410 420 430
AVVENVPPLRKEFVRRGLSDHEIDRLELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRDMDDLG
||||||||||||||||||||||||||||||||||||||||||||||||||||||||
AVVENVPPLRKEFVRRGLSDHEIDRLELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRDMDDLG
370 380 390 400 410 420 430

440 450 X
CLEDIEEALCGPAALPPAPSLLR
|||||||||||||||||||
CLEDIEEALCGPAALPPAPSLLR
440 450 X

4. US-07-625-668A-2 (1-455)

R07451 Human Tumour Necrosis Factor-Receptor from lambdaT

ID R07451 standard; protein; 455 AA.

AC R07451;

DT 29-JAN-1991 (first entry)

DE Human Tumour Necrosis Factor-Receptor from lambdaTNF-R2 cDNA insert.

KW Tumour necrosis factor binding protein; TNF-B; TNF-receptor;

KW infectious disease; parasitic disease; cachexia;

KW autoimmune disease; shock; lambdaTNF-R2; ratNF-R8.

OS Homo sapiens.
PN EP-393438-A.
PD 24-OCT-1990.
PF 06-APR-1990; 106624.
PR 21-APR-1989; DE-913101.
PR 21-JUN-1989; DE-920282.
PA (BOEH) BOEHRINGER INGELHEIMINT.
PI Hauptmann R, Himmller A, Maurer-Fogy I, Stratowa C;
DR WPI; 90-321987/43.
DR N-PSDB; Q06285.
PT DNA encoding TNF binding protein and TNF- receptor - used in
PT tumour treatment and to understand mechanismsm to TNF action
PS Disclosure; Fig 91(1-2); 51pp; German.
CC ratTNF-R8 (Q06284) was used to screen the HS913T cDNA library.
CC LambdaTNF-R2 encodes the complete human TNF-R2 and was used to
CC construct a plasmid (pADTNF-R) expressing the product the same way
CC as pADTNF-BP (see Q06282). The expressed proteins are useful
CC prophylactically and therapeutically to control disorders which
CC involve the damaging effects of TNF-alpha or -beta (e.g. infectious or
CC parasitic diseases, shock, cachexia, autoimmune diseases, adult
CC respiratory distress syndrome etc., or side effects of treatment with
CC TNF-alpha). They can also be used as diagnostic reagents for
CC assaying TNF and in study of TNF-receptor interactions.
CC See also Q06282-Q06285.
SQ Sequence 455 AA;
SQ 21 A; 27 R; 17 N; 21 D; 0 B; 30 C; 16 Q; 29 E; 0 Z; 29 G; 10 H;
SQ 13 I; 57 L; 20 K; 5 M; 14 F; 37 P; 37 S; 30 T; 5 W; 13 Y; 24 V;

Initial Score = 451 Optimized Score = 451 Significance = 40.97
Residue Identity = 99% Matches = 451 Mismatches = 4
Gaps = 0 Conservative Substitutions = 0

X 10 20 30 40 50 60 70
MGLSTVPDLLLPLVLELLVGIYPSGVIGLVPHLGDREKRDSVCPQGKYIHPQNNNSICCTKCHKGTYLYND
||||| ||||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
MGLSSVPDLLLPLVLELLVGIYPSAVIGLVPHLGDREKRDSVCPQGKYIHPQNNNSICCTKCHKGTYLYND
X 10 20 30 40 50 60 70

80 90 100 110 120 130 140
PGPGQDTDCRECESGSFTASENHRLHCLSCSKCRKEMGQVEISSCTVDRDTVCGRKNGYRHYWSENLFQCF
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
PGPGQDTDCRECESGSFTASENHRLHCLSCSKCRKEMGQVEISSCTVDRDTVCGRKNGYRHYWSENLFQCF
80 90 100 110 120 130 140

150 160 170 180 190 200 210
NCSLCLNGTVHLSCQEKKQNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
NCSLCLNGTVHLSCQEKKQNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL
150 160 170 180 190 200 210

220 230 240 250 260 270 280
VIFFGLCLLSLLFIGLMYRYQRWKSCLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFS
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
VIFFGLCLLSLLFIGLMYRYQRWKSCLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFS
220 230 240 250 260 270 280

290 300 310 320 330 340 350 360
PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQGADPILATALASDPIPPLQKWEDEAHKPQSLDDPATLY
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQGADPILATALASDPIPPLQKWEDEAHKPQSLDDPATLY
290 300 310 320 330 340 350 360

370 380 390 400 410 420 430
AVVENVPPLRWKEFVRLGLSDHEIDRLELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRDMDDLG
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
AVVENVPPLRWKEFVRLGLSDHEIDRLELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRDMDDLG

220	230	240	250	260	270	280
VIFFGLCLLSLLFIGLMYR	Y	W	Y	Y	Y	Y
VIFFGLCLLSLLFIGLMYR	Y	W	Y	Y	Y	Y
220	230	240	250	260	270	280

290	300	310	320	330	340	350	360
PSSTFTSSSTYTPGDCPNFA	APPREVAPPY	QGADPILATALASD	PIPNPLQKWE	DSAHKPQS	LDTDDP	ATLY	
PSSTFTSSSTYTPGDCPNFA	APPREVAPPY	QGADPILATALASD	PIPNPLQKWE	DSAHKPQS	LDTDDP	ATLY	
290	300	310	320	330	340	350	360

X	380
AVVENVPPLRWKEFVRRRLGLS	
AVVENVPPLRW	
370	

6. US-07-625-668A-2 (1-455)

R07450 Rat Tumour Necrosis Factor-Receptor from raTNF-R8

ID R07450 standard; protein: 461 AA.
 AC R07450;
 DT 29-JAN-1991 (first entry)
 DE Rat Tumour Necrosis Factor-Receptor from raTNF-R8 cDNA.
 KW Tumour necrosis factor binding protein; TNF-BP; TNF-receptor;
 KW raTNF-R8.
 OS Rat *rattus*.
 PN EP-393438-A.
 PD 24-OCT-1990.
 PF 06-APR-1990; 106624.
 PR 21-APR-1989; DE-913101.
 PR 21-JUN-1989; DE-920282.
 PA (BOEH) BOEHRINGER INGELHEIMINT.
 PI Hauptmann R, Himmller A, Maurer-Fogy I, Stratowa C;
 DR WPI; 90-321987/43.
 DR N-PSDB; Q06284.
 PT DNA encoding TNF binding protein and TNF- receptor - used in
 PT tumour treatment and to understand mechanismsm to TNF action
 PS Disclosure; Fig 8(1-2); 51pp; German.
 CC A rat brain cDNA analogue of the HS913T cDNA library from rat
 CC glioma cell line C6 (ATCC CCL107) is prepared in lambda-gt11.
 CC The isolated clone raTNF-R8 is used as probe to isolated the entire
 CC human TNF receptor, as represented in Q06285.
 CC See also Q06282-Q06285.
 SQ Sequence 461 AA;
 SQ 22 A; 26 R; 20 N; 16 D; 0 B; 31 C; 16 Q; 25 E; 0 Z; 28 G; 14 H;
 SQ 16 I; 48 L; 20 K; 9 M; 17 F; 44 P; 31 S; 29 T; 5 W; 8 Y; 36 V;

Initial Score = 240 Optimized Score = 212 Significance = 19.02
 Residue Identity = 48% Matches = 223 Mismatches = 232
 Gaps = 6 Conservative Substitutions = 0

X	10	20	30	40	50	60	70
MGLSTVPDLLLLPLVLLELLVGIYPSGVIGLVPHLGDREKRDSVCPQGKYIHPQNNNSICCTKCHKGTYLYNDC							
MGLPIVPGLLLLSLVLLALLMGIHPGVTGLVPSLGDREKRDNLCPQGKYAHPKNNNSICCTKCHKGTYLVSDC							
X	10	20	30	40	50	60	70

80	90	100	110	120	130	140
PGPGQDTCRECESGSFTASENHLRHC	LCSKCRKEMGQVEISSCTVDRDTVCGCRKNQYRHYWSENLFQCF					
PGPGQETVCELSHKGFTASQNHVRQCL	CLSCKTCRKEMFQVEISPCKADM					
80	90	100	110	120	130	140

370

380

390

400

410

420

430

440 450 X
 CLEDIEEALCGPAALPPAPSLLR
 |||||||||||||||||||||
 CLEDIEEALCGPAALPPAPSLLR
 440 450 X

5. US-07-625-668A-2 (1-455)

R07449 Tumour Necrosis Factor-Binding Protein from pTNF-B

ID R07449 standard; protein; 371 AA.
 AC R07449;
 DT 29-JAN-1991 (first entry)
 DE Tumour Necrosis Factor-Binding Protein from pTNF-BP15 cDNA.
 KW Tumour necrosis factor binding protein; TNF-BP; TNF-receptor;
 KW pTNF-BP15; infectious disease; parasitic disease; cachexia;
 KW autoimmune disease; shock.
 OS Homo sapiens.
 PN EP-393438-A.
 PD 24-OCT-1990.
 PF 06-APR-1990; 106624.
 PR 21-APR-1989; DE-913101.
 PR 21-JUN-1989; DE-920282.
 PA (BOEH) BOEHRINGER INGELHEIMINT.
 PI Hauptmann R, Himmller A, Maurer-Fogy I, Stratowa C;
 DR WPI; 90-321987/43.
 DR N-PSDB; Q06282.
 PT DNA encoding TNF binding protein and TNF- receptor - used in
 PT tumour treatment and to understand mechanismsm to TNF action
 PS Disclosure; Fig 1(1-3); 51pp; German.
 CC Clone pTNF-BP15 was used to contruct pADTNF-BP, for transfection of
 CC e.g. COS7 cells. The expressed proteins are useful
 CC prophylactically and therapeutically to control disorders which
 CC involve the damaging effects of TNF-alpha or -beta (e.g. infectious or
 CC parasitic diseases, shock, cachexia, autoimmune diseases, adult
 CC respiratory distress syndrome etc., or side effects of treatment with
 CC TNG-alpha). They can also be used as diagnostic reagents for
 CC assaying TNF and in study of TNF-receptor interactions.
 CC See also Q06282-Q06285.
 SQ Sequence 371 AA;
 SQ 13 A; 14 R; 16 N; 16 D; 0 B; 27 C; 13 Q; 21 E; 0 Z; 25 G; 9 H;
 SQ 11 I; 41 L; 19 K; 3 M; 13 F; 32 P; 33 S; 28 T; 4 W; 12 Y; 21 V;

Initial Score = 371 Optimized Score = 371 Significance = 33.62
 Residue Identity = 100% Matches = 371 Mismatches = 0
 Gaps = 0 Conservative Substitutions = 0

X 10 20 30 40 50 60 70
 MGLSTVPDLLLPLVLLELLVGIYPSGVIGLVPHLGDREKRDSVCPQGKYIHPQNNSICCTKCHKGTYLYNDC

||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

MGLSTVPDLLLPLVLLELLVGIYPSGVIGLVPHLGDREKRDSVCPQGKYIHPQNNSICCTKCHKGTYLYNDC

X 10 20 30 40 50 60 70

80 90 100 110 120 130 140
 PGPGQDTDCRECESGSFTASENHLRHCLSCSKCRKEMGQVEISSCTVDRDTVCGCRKNQYRHYWSENLFQCF

||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

PGPGQDTDCRECESGSFTASENHLRHCLSCSKCRKEMGQVEISSCTVDRDTVCGCRKNQYRHYWSENLFQCF

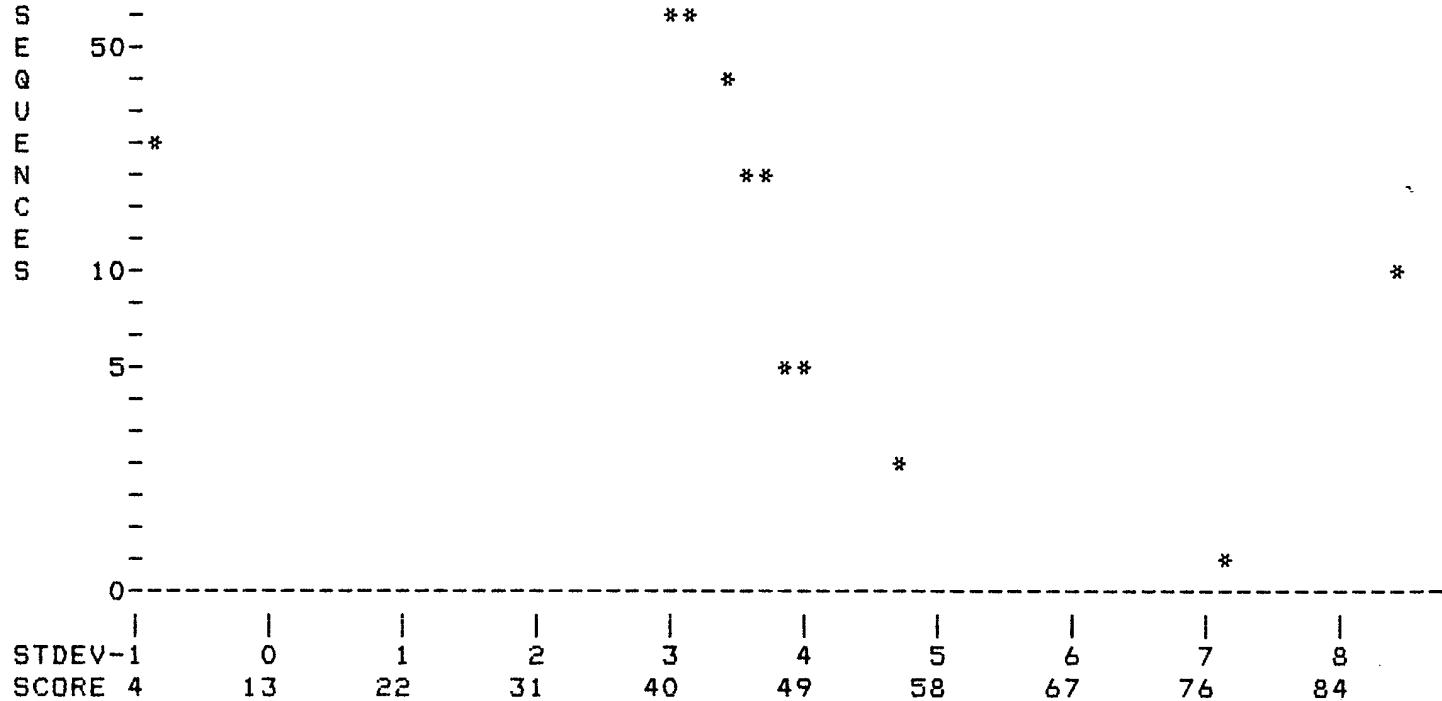
80 90 100 110 120 130 140

150 160 170 180 190 200 210
 NCSLCLNGTVHLSCQEKGNTVCTCHAGFFLRENECVSCSNCKSLECTKLCCLPQIENVKGTEDSGTTVLLPL

||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

NCSLCLNGTVHLSCQEKGNTVCTCHAGFFLRENECVSCSNCKSLECTKLCCLPQIENVKGTEDSGTTVLLPL

150 160 170 180 190 200 210



PARAMETERS

Similarity matrix	Unitary	K-tuple	2
Mismatch penalty	5	Joining penalty	20
Gap penalty	5.00	Window size	13
Gap size penalty	0.26		
Cutoff score	0		
Randomization group	0		
Initial scores to save	20	Alignments to save	10
Optimized scores to save	20	Display context	10

SEARCH STATISTICS

Scores:	Mean	Median	Standard Deviation
	13	25	9.09
Times:	CPU	Total Elapsed	
	00:02:47.05	00:05:40.00	
Number of residues:	1849227		
Number of sequences optimized:	4342		

The scores below are sorted by optimized score.
 Significance is calculated based on optimized score.

4 100% identical sequences to the query sequence were found:

Sequence Name	Description	Init. Opt.				
		Length	Score	Score	Sig.	Frame
1. A34899	*Tumor necrosis factor recept	455	455	455	48.61	0
2. A34900	*Tumor necrosis factor recept	455	455	455	48.61	0
3. S12057	*Tumor necrosis factor recept	455	455	455	48.61	0
4. A36555	*Tumor necrosis factor recept	455	455	455	48.61	0

The list of other best scores is:

Sequence Name	Description	Init. Opt.				
		Length	Score	Score	Sig.	Frame

Number of scores above cutoff:

4342

Cut-off raised to 4.
Cut-off raised to 5.
Cut-off raised to 6.
Cut-off raised to 7.

The scores below are sorted by initial score.
Significance is calculated based on initial score.

4 100% identical sequences to the query sequence were found:

Sequence Name	Description	Init. Opt.				
		Length	Score	Score	Sig.	Frame
1. A34899	*Tumor necrosis factor recept	455	455	455	97.80	0
2. A34900	*Tumor necrosis factor recept	455	455	455	97.80	0
3. S12057	*Tumor necrosis factor recept	455	455	455	97.80	0
4. A36555	*Tumor necrosis factor recept	455	455	455	97.80	0

The list of other best scores is:

Sequence Name	Description	Init. Opt.				
		Length	Score	Score	Sig.	Frame
	**** 97 standard deviations above mean ****					
5. A38281	*Tumor necrosis factor recept	455	452	452	97.15	0
	**** 51 standard deviations above mean ****					
6. B36555	*Tumor necrosis factor recept	461	243	215	51.83	0
	**** 47 standard deviations above mean ****					
7. B40254	*Tumor necrosis factor recept	454	224	267	47.71	0
8. A38634	*Tumor necrosis factor recept	454	224	267	47.71	0
9. S16677	*P55 tumor necrosis factor re	454	224	267	47.71	0
	**** 11 standard deviations above mean ****					
10. C36555	*Tumor necrosis factor bindin	134	55	78	11.06	0
	**** 3 standard deviations above mean ****					
11. A38258	*Tumor necrosis factor blocki	20	20	20	3.47	0
	**** 1 standard deviation above mean ****					
12. A42502	*C22L protein - Vaccinia viru	122	11	21	1.52	0
13. I42528	*B28R protein - Vaccinia viru	122	11	21	1.52	0
14. W5WL11	E5A protein - Human papilloma	91	10	16	1.30	0
15. S00897	Pancreatic ribonuclease A pre	150	10	24	1.30	0
16. A31556	Glucose-transporter protein,	522	10	30	1.30	0
17. S05319	Glucose transport protein, he	523	10	28	1.30	0
18. S06920	Glucose transport protein, he	523	10	28	1.30	0
19. S01085	Hypothetical protein 1 - Whit	1294	10	43	1.30	0
20. A35186	*Salivary agglutinin receptor	1473	10	39	1.30	0

Query sequence being compared: US-07-625-668A-2 (1-455)

Number of sequences optimized: 4342

Results of the optimized comparison of US-07-625-668A-2 (1-455) with:
Data bank : PIR 32, all entries

1000-
-
N -
U 500-
M - * *
B -
E - * * * *
R - ** ** * *
- * * * *
O - ** **
F 100- * ** *
- * ** *

**** 48 standard deviations above mean ****
5. A38281 *Tumor necrosis factor recept 455 452 452 48.28 0
**** 7 standard deviations above mean ****
6. S16677 *P55 tumor necrosis factor re 454 224 267 27.93 0
7. B40254 *Tumor necrosis factor recept 454 224 267 27.93 0
8. A38634 *Tumor necrosis factor recept 454 224 267 27.93 0
**** 22 standard deviations above mean ****
9. B36555 *Tumor necrosis factor recept 461 243 215 22.21 0
**** 7 standard deviations above mean ****
10. C36555 *Tumor necrosis factor bindin 134 55 78 7.15 0
**** 4 standard deviations above mean ****
11. B38634 *Tumor necrosis factor recept 474 8 55 4.62 0
12. A40254 *Tumor necrosis factor recept 474 8 55 4.62 0
**** 3 standard deviations above mean ****
13. S18984 *Arrestin - Human (fragment) 409 8 49 3.96 0
14. GQHUN Nerve growth factor receptor 427 9 49 3.96 0
15. OYB077 Guanylate cyclase, soluble, 7 691 6 49 3.96 0
16. S15921 *Protein TPX-VT3 - Thermoprot 474 8 49 3.96 0
17. B24785 Hypothetical protein 1028 - S 1028 6 48 3.85 0
18. A26728 Elastin a precursor - Bovine 747 7 48 3.85 0
19. A36226 *Collagen alpha-1 - Sea urchi 730 7 48 3.85 0
20. A26456 Nicotinic acetylcholine recep 625 7 48 3.85 0

1. US-07-625-668A-2 (1-455)

A34899 *Tumor necrosis factor receptor precursor, 55K -

ENTRY A34899 #Type Protein
TITLE *Tumor necrosis factor receptor precursor, 55K -
Human
DATE 01-Aug-1990 #Sequence 01-Aug-1990 #Text 01-Aug-1990
PLACEMENT 0.0 0.0 0.0 0.0 0.0
COMMENT *This entry is not verified.
REFERENCE
#Authors Loetscher H., Pan Y.C.E., Lahm H.W., Gentz R.,
Brockhaus M., Tabuchi H., Lesslauer W.
#Journal Cell (1990) 61:351-359
#Title Molecular cloning and expression of the human 55 kd
tumor necrosis factor receptor.
#Reference-number A34899
#Accession A34899
SUMMARY #Molecular-weight 50494 #Length 455 #Checksum 153
SEQUENCE

Initial Score = 455 Optimized Score = 455 Significance = 48.61
Residue Identity = 100% Matches = 455 Mismatches = 0
Gaps = 0 Conservative Substitutions = 0

X 10 20 30 40 50 60 70
MGLSTVPDLLLLPLVLLELLVGIYPSGVIGLVPHLGDREKRDSVCPQGKYIHPQNNSICCTKCHKGTYLYNDC
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
MGLSTVPDLLLLPLVLLELLVGIYPSGVIGLVPHLGDREKRDSVCPQGKYIHPQNNSICCTKCHKGTYLYNDC
X 10 20 30 40 50 60 70

80 90 100 110 120 130 140
PGPGQDTDCRECESGSFTASENHRLHCLSCSKCRKEMGQVEISSCTVDRDTVCGCRKNQYRHYSENLFQCF
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
PGPGQDTDCRECESGSFTASENHRLHCLSCSKCRKEMGQVEISSCTVDRDTVCGCRKNQYRHYSENLFQCF
80 90 100 110 120 130 140

150 160 170 180 190 200 210
NCSLCLNGTVHLSCQEKGNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
NCSLCLNGTVHLSCQEKGNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL
150 160 170 180 190 200 210

220	230	240	250	260	270	280	
VIFFGLCLLSLLFIGLMYRYQRWKSCLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFSPV							
VIFFGLCLLSLLFIGLMYRYQRWKSCLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFSPV							
220	230	240	250	260	270	280	
290	300	310	320	330	340	350	360
PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQGADPILATALASDPIPNNPLQKWEDEAHKPQSLDDPATLY							
PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQGADPILATALASDPIPNNPLQKWEDEAHKPQSLDDPATLY							
290	300	310	320	330	340	350	360
370	380	390	400	410	420	430	
AVVENVPPLRWKEFVRRGLSDHEIDRLELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRDM DLLG							
AVVENVPPLRWKEFVRRGLSDHEIDRLELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRDM DLLG							
370	380	390	400	410	420	430	
440	450	X					
CLEDIEEALCGPAALPPAPSLLR							
CLEDIEEALCGPAALPPAPSLLR							
440	450	X					

2. US-07-625-668A-2 (1-455)

A34900 *Tumor necrosis factor receptor precursor - Human

ENTRY A34900 #Type Protein
TITLE *Tumor necrosis factor receptor precursor - Human
DATE 01-Aug-1990 #Sequence 01-Aug-1990 #Text 01-Aug-1990
PLACEMENT 0.0 0.0 0.0 0.0 0.0
COMMENT *This entry is not verified.
REFERENCE
 #Authors Schall T.J., Lewis M., Koller K.J., Lee A., Rice
 G.C., Wong G.H.W., Gatanaga T., Granger G.A.,
 Lentz R., Raab H., Kohr W.J., Goeddel D.V.
 #Journal Cell (1990) 61:361-370
 #Title Molecular cloning and expression of a receptor for
 human tumor necrosis factor.
 #Reference-number A34900
 #Accession A34900
SUMMARY #Molecular-weight 50494 #Length 455 #Checksum 153
SEQUENCE

220	230	240	250	260	270	280	
VIFFGLCLLSLLFIGLMYRY	WKS	KLYSIVCGKSTPEKEGELEGTT		LAPNPSFSPTPGFTPTLGFS			
VIFFGLCLLSLLFIGLMYRYQRWKS	KLYSIVCGKSTPEKEGELEGTT	KPL	APNPSFSPTPGFTPTLGFS				
220	230	240	250	260	270	280	
290	300	310	320	330	340	350	360
PSSTFTSSSTYTPGDCPNFAAPR	REVAPPYQGADP	I	ALASD	PIPNPLQK	WEDSAHKP	QSLDTDD	PATLY
PSSTFTSSSTYTPGDCPNFAAPR	REVAPPYQGADP	I	ALASD	PIPNPLQK	WEDSAHKP	QSLDTDD	PATLY
290	300	310	320	330	340	350	360
370	380	390	400	410	420	430	
AVVENVPLRWKEFVRR	GLG	SDHEIDR	LELQNGRCL	REAQYSML	ATWRRRT	PRREAT	LELLGRV
AVVENVPLRWKEFVRR	GLG	SDHEIDR	LELQNGRCL	REAQYSML	ATWRRRT	PRREAT	LELLGRV
370	380	390	400	410	420	430	
440	450	X					
CLEDIEEALCGPAALPPAPSLLR							
CLEDIEEALCGPAALPPAPSLLR							
440	450	X					

3. US-07-625-668A-2 (1-455)

S12057 *Tumor necrosis factor receptor type 1 - Human?

ENTRY	S12057	#Type Protein			
TITLE	*Tumor necrosis factor receptor type 1 - Human				
DATE	09-Jul-1991 #Sequence 09-Jul-1991 #Text 09-Jul-1991				
PLACEMENT	0.0	0.0	0.0	0.0	0.0
COMMENT	*This entry is not verified.				
SOURCE	Homo sapiens #Common-name man				
REFERENCE	#Authors Nophar Y., Kemper O., Brakebusch C., Engelmann H., Zwang R., Aderka D., Holtmann H., Wallach D. #Journal EMBO J. (1990) 9:3269-3278 #Title Soluble forms of tumor necrosis factor receptors (TNF-Rs). The cDNA for the type I TNF-R, cloned using amino acid sequence data of its soluble form, encodes both the cell surface and a soluble form of the receptor. #Reference-number S12057 #Accession S12057				
SUMMARY	#Molecular-weight 50494 #Length 455 #Checksum 153				
SEQUENCE					

Initial Score = 455 Optimized Score = 455 Significance = 48.61
 Residue Identity = 100% Matches = 455 Mismatches = 0
 Gaps = 0 Conservative Substitutions = 0

X	10	20	30	40	50	60	70
MGLSTVPD	LLLPLV	LLVGIYPSGV	IGLVPHLG	DREKRDSVCP	QGKYIHPQNN	SICCTKCHKG	TYLYNDC
MGLSTVPD	LLLPLV	LLVGIYPSGV	IGLVPHLG	DREKRDSVCP	QGKYIHPQNN	SICCTKCHKG	TYLYNDC
X	10	20	30	40	50	60	70

80	90	100	110	120	130	140	
PGPGQDTDCRECESGSFT	ASENH	LRHCL	CSKCR	KEMGQ	VEISSCTV	DRDTVC	GCRKNQYRHY
PGPGQDTDCRECESGSFT	ASENH	LRHCL	CSKCR	KEMGQ	VEISSCTV	DRDTVC	GCRKNQYRHY
80	90	100	110	120	130	140	

150	160	170	180	190	200	210
-----	-----	-----	-----	-----	-----	-----

NCSLCLNGTVHLSCQEKGQNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL
 |||||||
 NCSLCLNGTVHLSCQEKGQNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL
 150 160 170 180 190 200 210
 220 230 240 250 260 270 280
 VIFFGLCLLSLLFIGLMYRYQRWKSCLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFSV
 |||||||
 VIFFGLCLLSLLFIGLMYRYQRWKSCLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFSV
 220 230 240 250 260 270 280
 290 300 310 320 330 340 350 360
 PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQQGADPILATALASDPIPPLQKWEDEAHKPQSLDDPATLY
 |||||||
 PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQQGADPILATALASDPIPPLQKWEDEAHKPQSLDDPATLY
 290 300 310 320 330 340 350 360
 370 380 390 400 410 420 430
 AVVENVPPLRWKEFVRRRLGLSDHEIDRLELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRDM DLLG
 |||||||
 AVVENVPPLRWKEFVRRRLGLSDHEIDRLELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRDM DLLG
 370 380 390 400 410 420 430
 440 450 X
 CLEDIEEALCGPAALPPAPSLLR
 |||||||
 CLEDIEEALCGPAALPPAPSLLR
 440 450 X

4. US-07-625-668A-2 (1-455)

A36555 *Tumor necrosis factor receptor precursor - Human }

ENTRY A36555 #Type Protein
 TITLE *Tumor necrosis factor receptor precursor - Human
 DATE 15-Apr-1991 #Sequence 15-Apr-1991 #Text 15-Apr-1991
 PLACEMENT 0.0 0.0 0.0 0.0 0.0
 COMMENT *This entry is not verified.
 SOURCE Homo sapiens #Common-name man
 REFERENCE
 #Authors Himmler A., Maurer-Fogy I., Kroenke M., Scheurich
 P., Pfizenmaier K., Lantz M., Olsson I., Hauptmann
 R., Stratowa C., Adolf G.R.
 #Journal DNA Cell Biol. (1990) 9:705-715
 #Title Molecular cloning and expression of human and rat
 tumor necrosis factor receptor chain (p60) and its
 soluble derivative, tumor necrosis factor-binding
 protein.
 #Reference-number A36555
 #Accession A36555
 SUMMARY #Molecular-weight 50494 #Length 455 #Checksum 153
 SEQUENCE

Initial Score = 455 Optimized Score = 455 Significance = 48.61
 Residue Identity = 100% Matches = 455 Mismatches = 0
 Gaps = 0 Conservative Substitutions = 0

X 10 20 30 40 50 60 70
 MGLSTVPDLLLPLVLELLVGIYPSGVIGLVPHLGDREKRDSCVCPQGKYIHPQNNNSICCTKCHKGTYLYNDC
 |||||||
 MGLSTVPDLLLPLVLELLVGIYPSGVIGLVPHLGDREKRDSCVCPQGKYIHPQNNNSICCTKCHKGTYLYNDC
 X 10 20 30 40 50 60 70
 80 90 100 110 120 130 140
 PGPGQDTCRECESGSFTASENHLRHCLSCSKCRKEMGQVEISSCTVDRDTVCGRKNQYRHYWSENLFQCF
 |||||||
 PGPGQDTCRECESGSFTASENHLRHCLSCSKCRKEMGQVEISSCTVDRDTVCGRKNQYRHYWSENLFQCF

PGPGQDTDCRECESGSFTASENHRLHCLSCSKCRKEMGQVEISSCTVDRDTVCGRKNQYRHYSSENLFQCF

80 90 100 110 120 130 140

150 160 170 180 190 200 210

NCSLCLNGLTVHLSCQEKGQNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL

||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

NCSLCLNGLTVHLSCQEKGQNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL

150 160 170 180 190 200 210

220 230 240 250 260 270 280

VIFFGLCLLSSLFIGLMYRYQRWKSCLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFSPV

||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

VIFFGLCLLSSLFIGLMYRYQRWKSCLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFSPV

220 230 240 250 260 270 280

290 300 310 320 330 340 350 360

PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQGADPILATALASDPIPNPLQKWEDESAHKPQSLDDPATLY

||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQGADPILATALASDPIPNPLQKWEDESAHKPQSLDDPATLY

290 300 310 320 330 340 350 360

370 380 390 400 410 420 430

AVVENVPPLRKEFVRLGLSDHEIDRLELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRDM DLLG

||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

AVVENVPPLRKEFVRLGLSDHEIDRLELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRDM DLLG

370 380 390 400 410 420 430

440 450 X

CLEDIEEALCGPAALPPAPSLLR

||||||||||||||||||||

CLEDIEEALCGPAALPPAPSLLR

440 450 X

5. US-07-625-668A-2 (1-455)

A38281 *Tumor necrosis factor receptor precursor - Human

ENTRY A38281 #Type Protein
TITLE *Tumor necrosis factor receptor precursor - Human
DATE 30-May-1991 #Sequence 30-May-1991 #Text 30-May-1991
PLACEMENT 0.0 0.0 0.0 0.0 0.0
COMMENT *This entry is not verified.
SOURCE Homo sapiens #Common-name man
REFERENCE
#Authors Gray P.W., Barrett K., Chantry D., Turner M.,
Feldmann M.
#Journal Proc. Natl. Acad. Sci. U.S.A. (1990) 87:7380-7384
#Title Cloning of human tumor necrosis factor (TNF)
receptor cDNA and expression of recombinant
soluble TNF-binding protein.
#Reference-number A38281
#Accession A38281
#Cross-reference GB:M37764
SUMMARY #Molecular-weight 50393 #Length 455 #Checksum 376
SEQUENCE

Initial Score = 452 Optimized Score = 452 Significance = 48.28

Residue Identity = 99% Matches = 452 Mismatches = 3

Gaps = 0 Conservative Substitutions = 0

X 10 20 30 40 50 60 70

MGLSTVPDLLLPLVLLVGIYPSGVIGLVPHLGDREKRDSVCPQGKYIHPQNNNSICCTKCHKGTYLYNDC

||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

MGLSTVPDLLLPLVLLVGIYPSGVIGLVPHLGDREKRDSVCPQGKYIHPQNNNSICCTKCHKGTYLYNDC

X 10 20 30 40 50 60 70

80	90	100	110	120	130	140		
PGPGQDTDCRECESGSFTASENHLRHCLSCSKCRKEMGQVEISSCTVDRDTVCGRKNQYRHYWSENLFQCF								
PGPGQDTDCRECESGSFTASENHLRHCLSCSKCRKEMGQVEISSCTVDRDTVCGRKNQYRHYWSENLFQCF	80	90	100	110	120	130	140	
150	160	170	180	190	200	210		
NCSLCLNGTVHLSCQEKGNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL								
NCSLCLNGTVHLSCQEKGNTVCTCHAGFFLRENECVSCSNCKSLECTKLCLPQIENVKGTEDSGTTVLLPL	150	160	170	180	190	200	210	
220	230	240	250	260	270	280		
VIFFGLCLLSLLFIGLMYRYQRWKSCLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFSPV								
VIFFGLCLLSLLFIGLMYRYQRWKSCLYSIVCGKSTPEKEGELEGTTKPLAPNPSFSPTPGFTPTLGFSPV	220	230	240	250	260	270	280	
290	300	310	320	330	340	350	360	
PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQGADPILATALASDPIPNNPLQKWEDEAHKPQSLDDPATLY								
PSSTFTSSSTYTPGDCPNFAAPRREVAPPYQGADPILATALASDPIPNNPLQKWEDEAHKPQSLDDPATLY	290	300	310	320	330	340	350	360
370	380	390	400	410	420	430		
AVVENVPLRWKEFVRRGLSDHEIDRLELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRDM DLLG								
AVVENVPLRTLTFVRRGLSDHEIDRLELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRNMDLLG	370	380	390	400	410	420	430	
440	450	X						
CLEDIEEALCGPAALPPAPSLLR								
CLEDIEEALCGPAALPPAPSLLR	440	450	X					

6. US-07-625-668A-2 (1-455)

S16677 *P55 tumor necrosis factor receptor - Mouse

ENTRY S16677 #Type Protein
TITLE *p55 tumor necrosis factor receptor - Mouse
DATE 07-Apr-1992 #Sequence 07-Apr-1992 #Text 07-Apr-1992
PLACEMENT 0.0 0.0 0.0 0.0 0.0
COMMENT *This entry is not verified.
SOURCE Mus musculus #Common-name house mouse
REFERENCE
#Authors Barrett K., Taylor-Fishwick D.A., Cope A.P.,
Kissonerghis A.M., Gray P.W., Feldmann M., Foxwell
B.M.J.
#Journal Eur. J. Immunol. (1991) 21:1649-1656
#Title Cloning, expression and cross-linking analysis of
the murine p55 tumor necrosis factor receptor.
#Reference-number S16677
#Accession S16677
#Cross-reference EMBL:X59238
SUMMARY #Molecular-weight 50129 #Length 454 #Checksum 4839
SEQUENCE

Initial Score = 224 Optimized Score = 267 Significance = 27.93
Residue Identity = 62% Matches = 287 Mismatches = 164
Gaps = 6 Conservative Substitutions = 0

X 10 20 30 40 50 60 70
MGLSTVPDLLLPLVLELLVGIVYPSGVIGLVPHLGDRKRDSCVCPQGKVIHPQNNNSICCTKCHKGTYLYNDC
||||| ||||| ||||| ||| ||| ||||| ||||| ||||| ||||| ||| ||||| ||||| ||||| ||| |||